

Bonine I

Phylogenetics (ch 27)

Thursday
29 January 2009
ECOL 182R UofA
K. E. Bonine

Kevin Bonine, Ph.D.

Tucson ~native

University of [Arizona](#) (undergrad)

- Ecology & Evolutionary Biology
- Economics

University of Wisconsin, Madison (graduate)

Zoology, [Evolutionary Physiology](#)
[Reptiles & Amphibians](#)

Teaching at UA since 2002

Herpetology
Vertebrate Physiology
Conservation Biology
Environmental Biology
Introductory Biology

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M 1:10-2pm

W 11:10-11am

in BSE113

Mostly Middle-Third of this course (plants etc.)

Text readings are highly encouraged

Website: For my lecture material:

<http://www.eebweb.arizona.edu/courses/182-spring2009-Bonine/182-BONINE-sp2009.htm>

There is a **link from Dr. Schaffer's 182R website.**

Text: The text is Freeman, Scott. 2008. *Biological Science* (Third Edition). Pearson Benjamin Cummings, San Francisco, CA. Available at UA Bookstore.

Posted Lecture Notes

Items in **orange** will not be available on-line (except for today), but they will be presented during lecture.

Note that slides are numbered for easy reference.

I will strive to post the lecture PDF file on your D2L site **before** lecture.



How do we organize/categorize biodiversity? Why would we want to?



Linnaean Taxonomy (1700s)

- **Taxonomy** is the effort to name and classify organisms.
- In Linnaeus' taxonomic system for classifying organisms, each organism is given a unique two-part scientific name consisting of the **genus** and the **species**.
 - (1) A **genus** is made up of a closely related group of species.
 - (2) A **species** is made up of individuals that regularly breed together and/or have characteristics that are distinct from those of other species.

Taxonomic Levels

- Linnaeus' system is **hierarchical** with nested **taxa**. The taxonomic levels from least to most specific are as follows:

domain
kingdom
phylum
class
order
family
genus
species

Linnaean hierarchy

- Group of organisms treated as unit is a **taxon** (plural **taxa**)
- Hierarchy of taxonomic categories

Kingdom	Plantae	Less specific ↑	Kingdom	Animalia
± 275,000 species			> 1,500,000 species	
Phylum	Angiospermae		Phylum	Chordata
± 250,000 species			± 50,000 species	
Class	Eudicotyledonae		Class	Aves
± 235,000 species			8,600 species	
Order	Rosales		Order	Passeriformes
± 18,000 species		5,160 species		
Family	Rosaceae	Family	Parulidae	
± 3,500 species		125 species		
Genus	<i>Rosa</i>	Genus	<i>Dendroica</i>	
± 500 species		28 species		
Species	<i>Rosa gallica</i>	Species	<i>Dendroica fusca</i>	
French rose		Blackburnian warbler		
		More specific ↓		

Italics



Linnaeus' Taxonomic Levels

KINGDOM
(Animalia)



PHYLUM
(Chordata)



CLASS
(Mammalia)



ORDER
(Primates)



FAMILY
(Hominidae)



GENUS
(*Homo*)



SPECIES
(*Homo sapiens*)



Theory of common descent

- Any two organisms can trace back to a common ancestor
- We all belong to a big family tree, some more closely related than others
- On the right is a history of **individuals**: can also draw up the history of **species**

(President) George
Walker Bush
b. 1946



(Senator) John
Forbes Kerry
b. 1943

Phylogeny

- **Phylogeny** = history of exactly how a group of organisms are descended from their common ancestor
- **Phylogenetic tree** = representation of that history

Lamarck

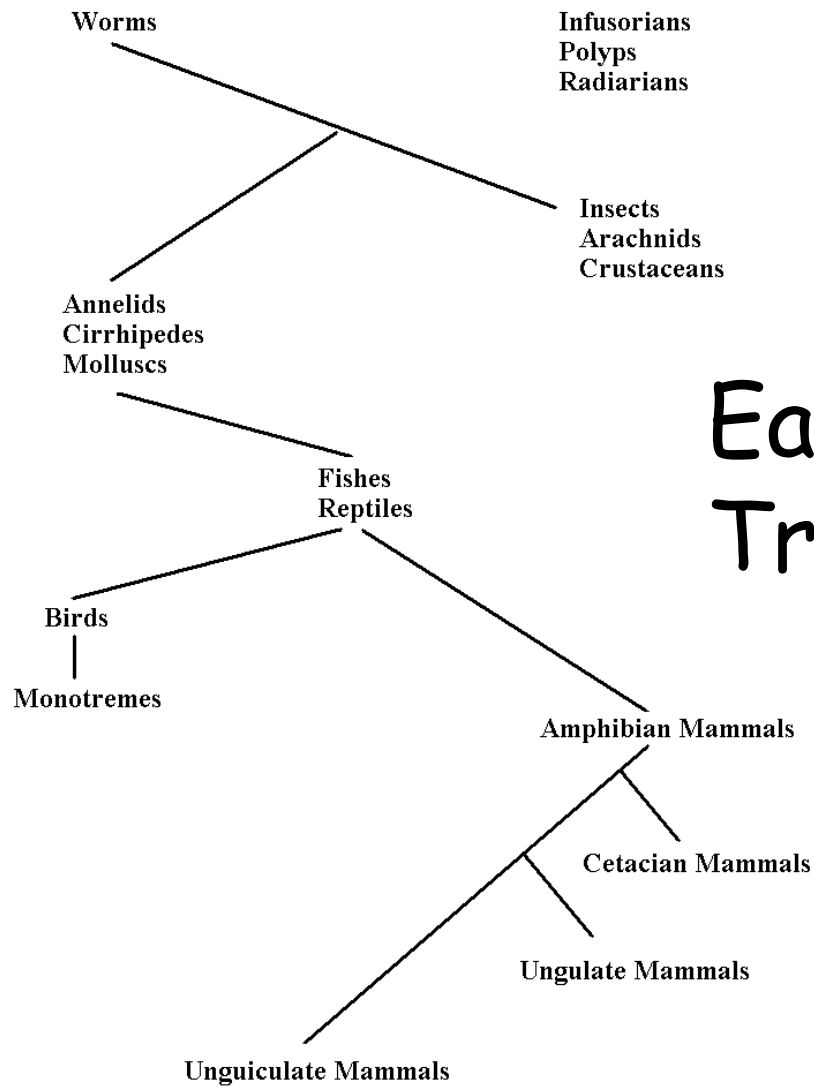


1821. From
Desmond 1989,
p. 43.

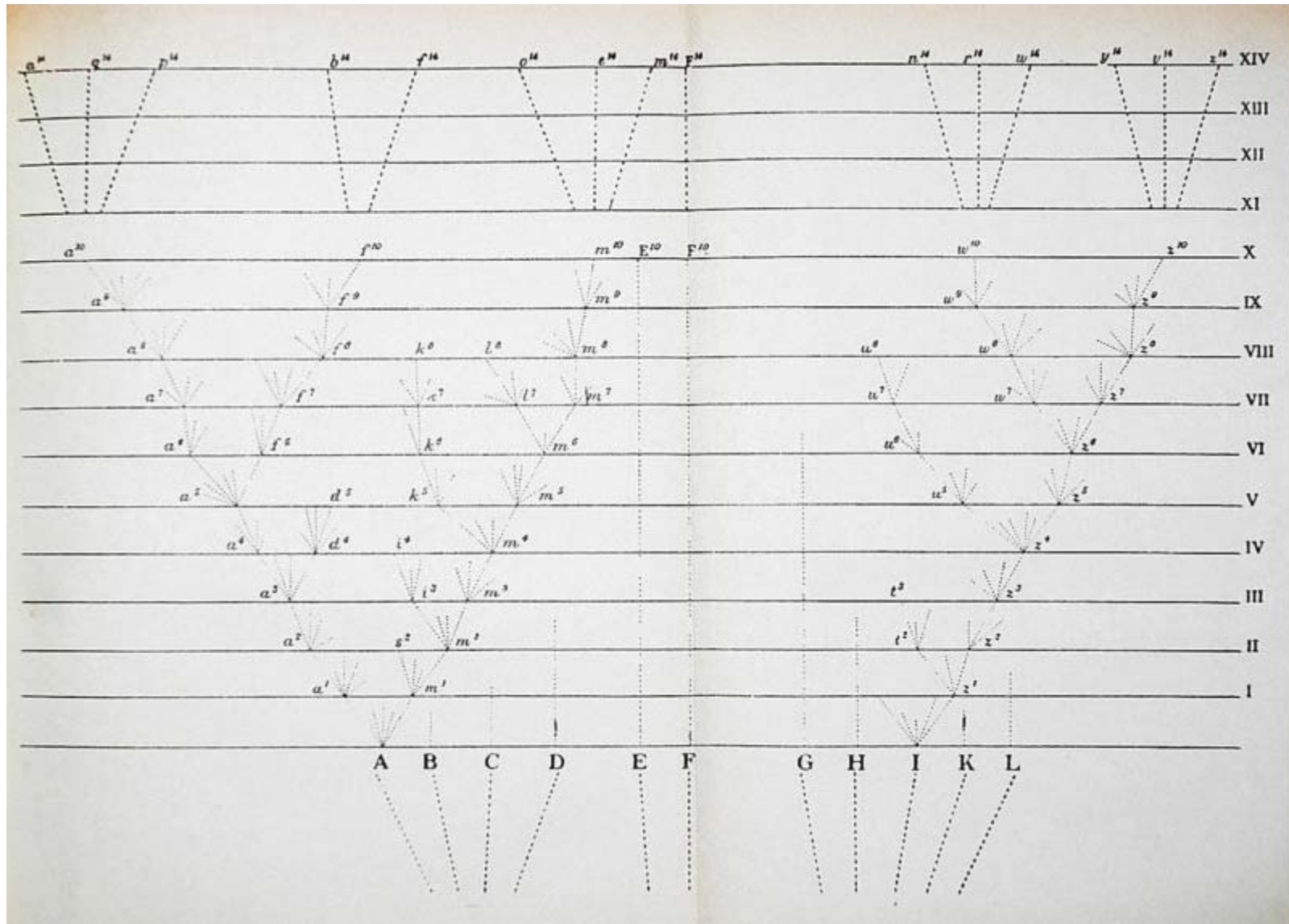


"Cineraria"

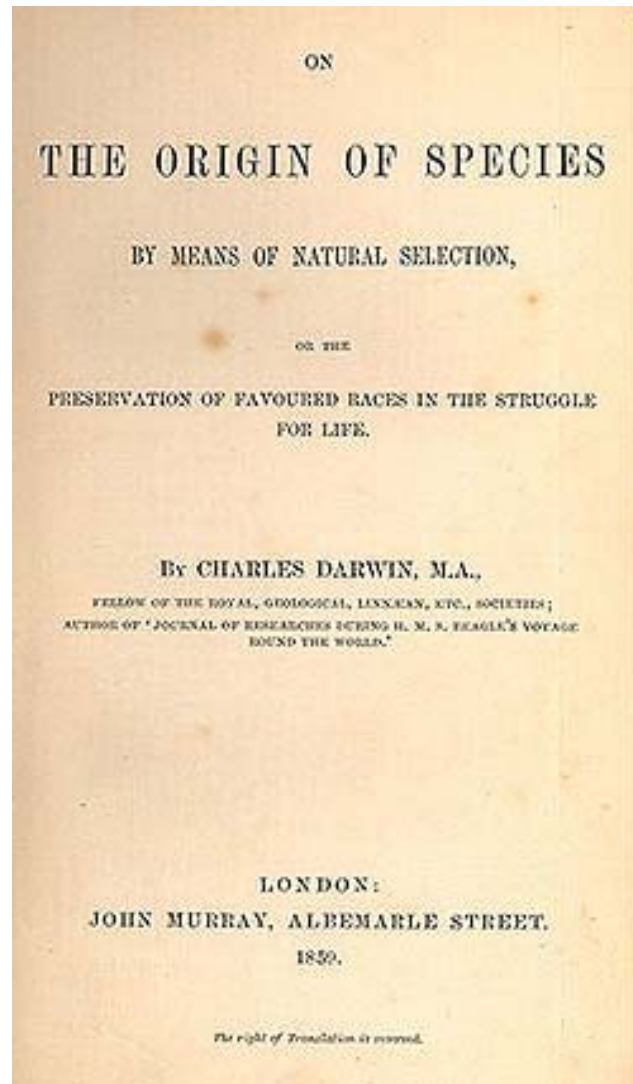
Jean Baptiste Pierre Antoine de Monet de Lamarck¹²
(1744-1829): *Tableau encyclopedique et methodique des
trois regnes de la nature... botanique*. Paris, 1791-1823



Early Phylogenetic Tree by Lamarck



Tree diagram used to show the divergence of species. It is the only illustration in *The Origin of Species* – Darwin 1859.

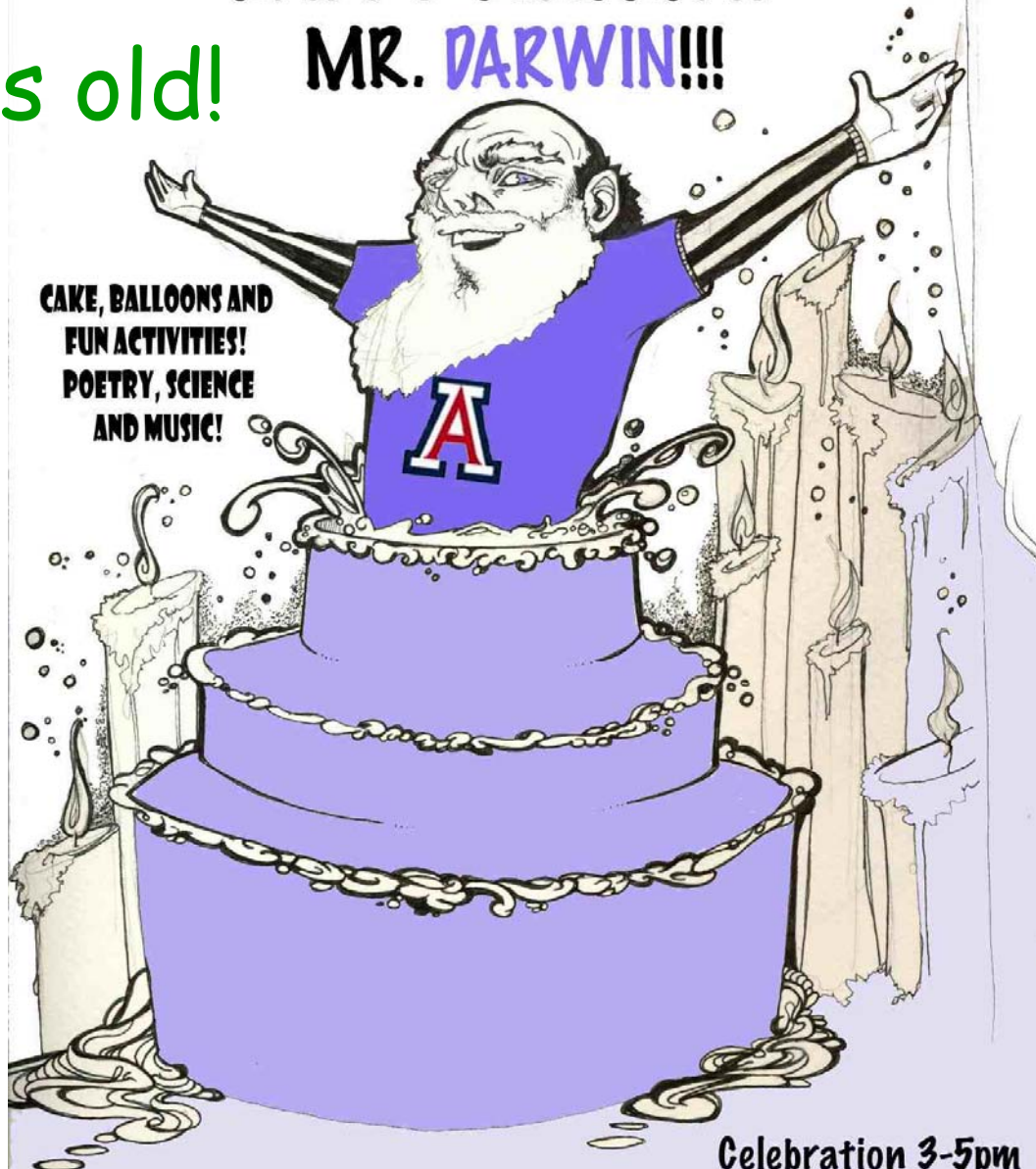


Title page of the 1859 edition
of *On the Origin of Species*

200 years old!

HAPPY BIRTHDAY MR. DARWIN!!!

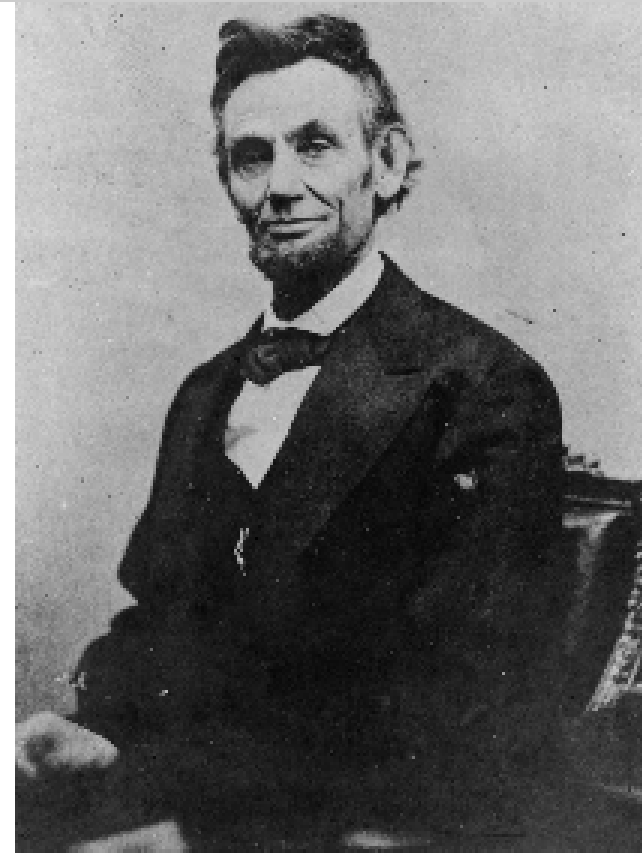
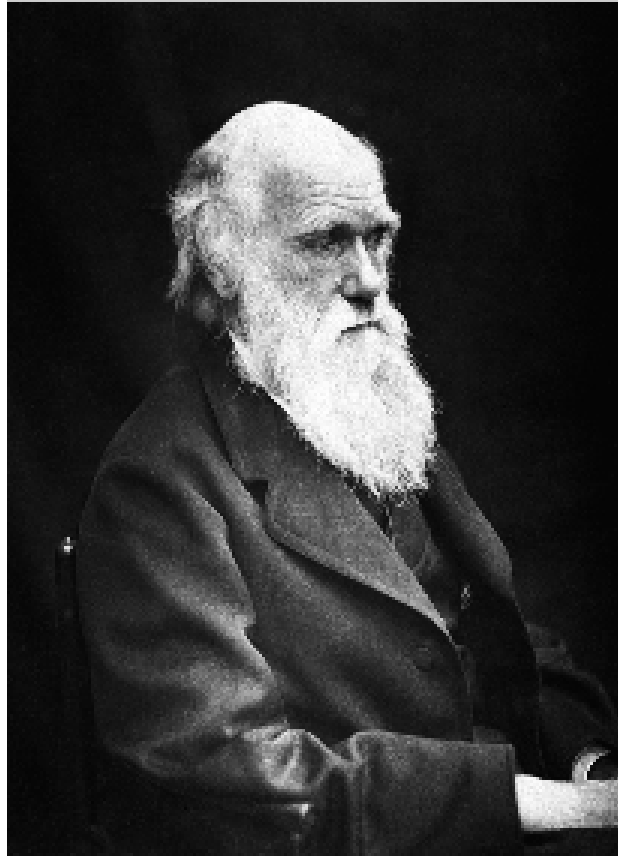
**CAKE, BALLOONS AND
FUN ACTIVITIES!
POETRY, SCIENCE
AND MUSIC!**



12 Feb
2009

sponsored by EEB, Poetry Center,
the UA Bookstore & others at UA

**Celebration 3-5pm
Thursday Feb 12
Student Union
Memorial Ballroom**

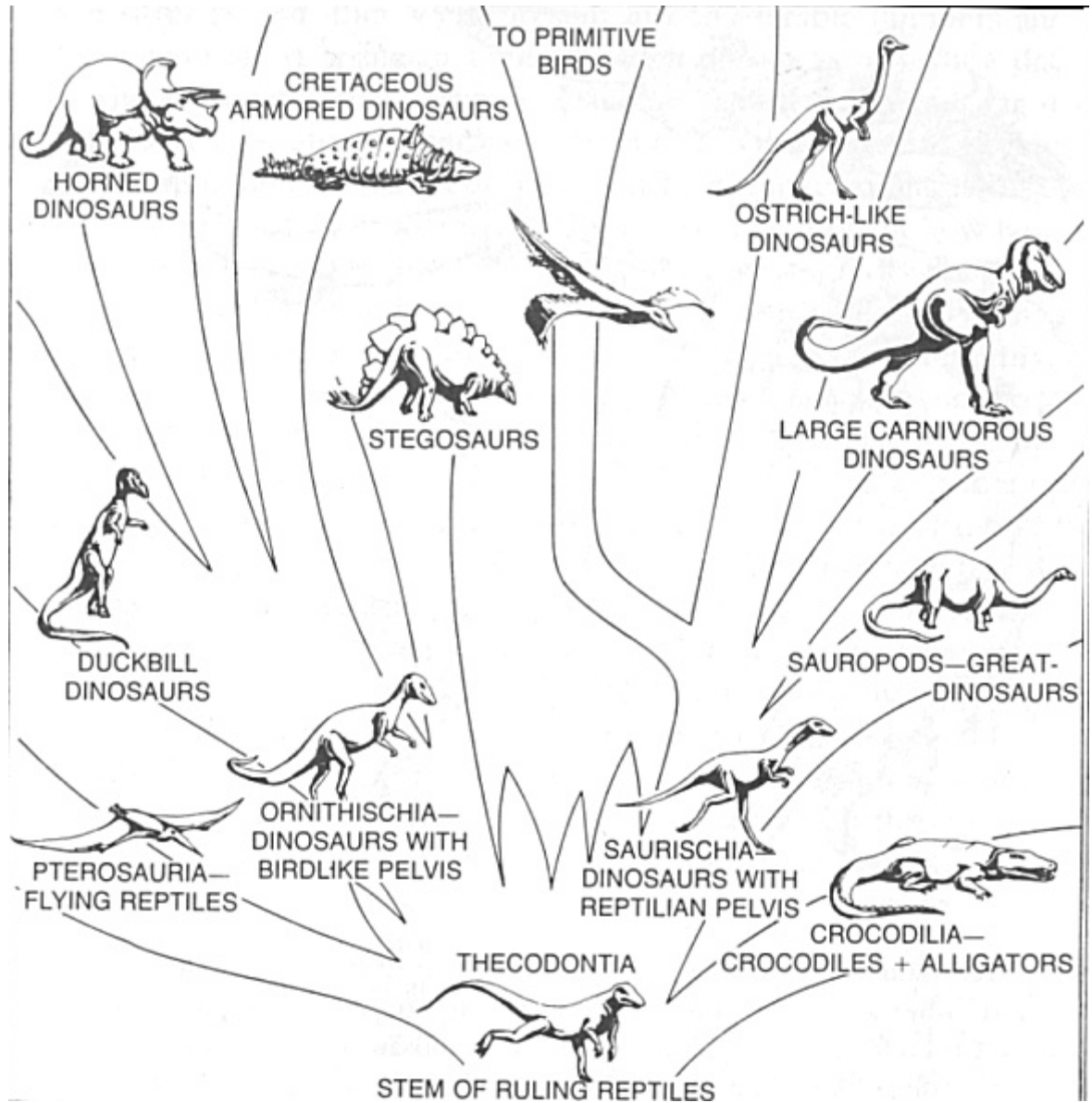


They Still Free Minds and Peoples
Darwin & Lincoln's Birthday, February 12th, 1809

- A **phylogenetic tree** is a graphical representation of the evolutionary relationships among species. Phylogenies can be established by analyzing similarities and differences in **traits**.



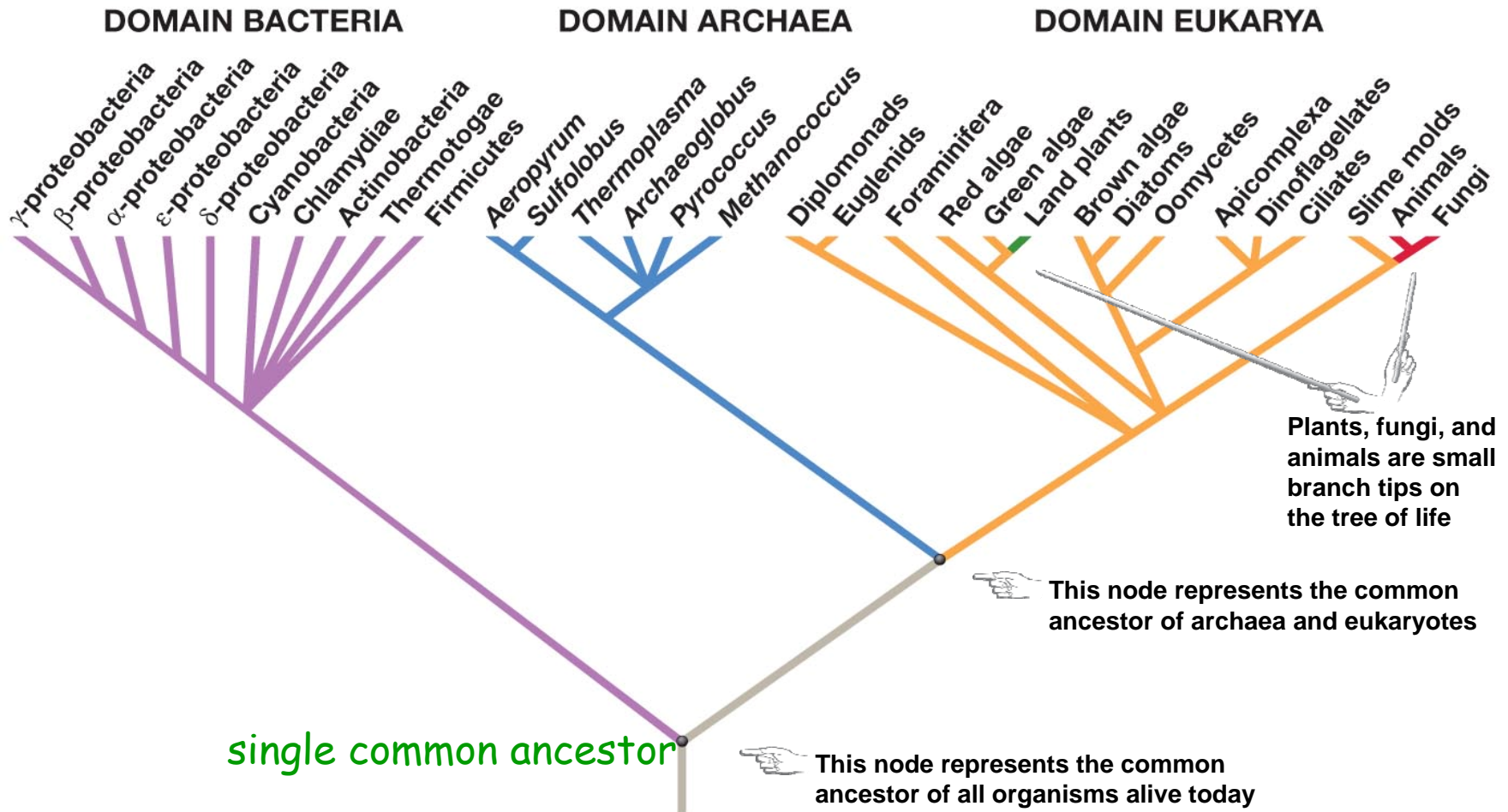
Often we are attempting to identify relationships from the distant **past**...



Phylogenetic Trees

- **Systematics**, the study of organismal diversity with respect to evolutionary (or not) relationships between organisms (patterns of descent).
 - **Taxonomy** - a subdiscipline that relates to classification
- What evolutionary relationships could be useful/helpful to understand?

Phylogenetic Tree of Life



Remember that >99% of all species are EXTINCT!

Old species often become new species...

Darwin's eventual conclusions stemming from his first question about the birds and plants of the Galapagos were to feature in one of the most important passages in *Origin of species* (pp. 397-406). The passage ended with one of his key points about evolution by natural selection:

The relations just discussed ... [including] the very close relation of the **distinct species** which inhabit the islets of the same archipelago, and especially the striking relation of the inhabitants of each whole archipelago or island to those of the nearest mainland, are, I think, **utterly inexplicable on the ordinary view of the independent creation of each species**, but are explicable on the view of **colonisation** from the nearest and readiest source, together with the **subsequent modification and better adaptation of the colonists to their new homes**.

http://darwin-online.org.uk/EditorialIntroductions/Chancellor_Keynes_Galapagos.html

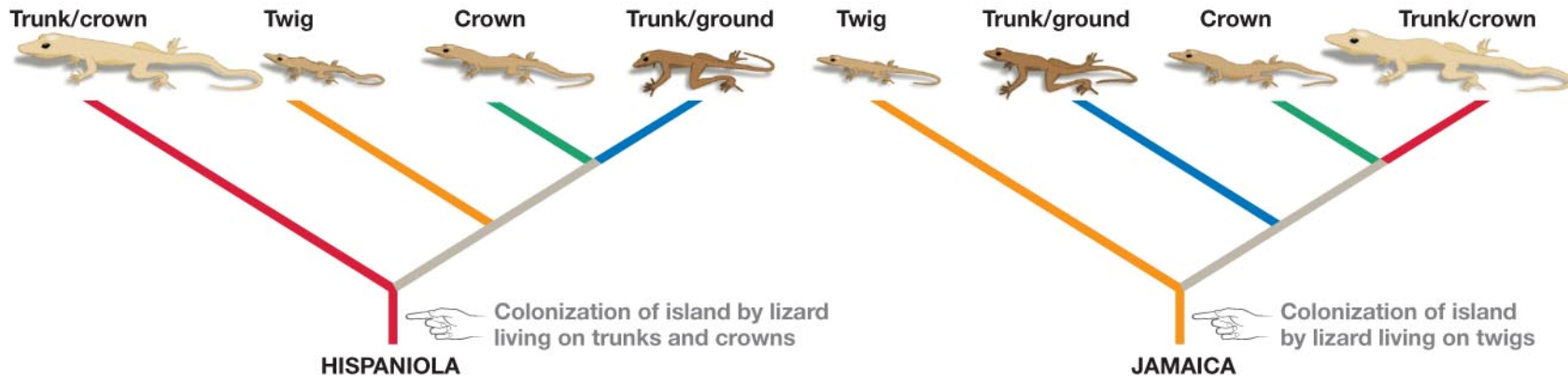


Adaptive Radiation

(a) Species of *Anolis* vary in leg length and tail length. Some species are ground dwelling; others live in distinct regions of shrubs or trees.

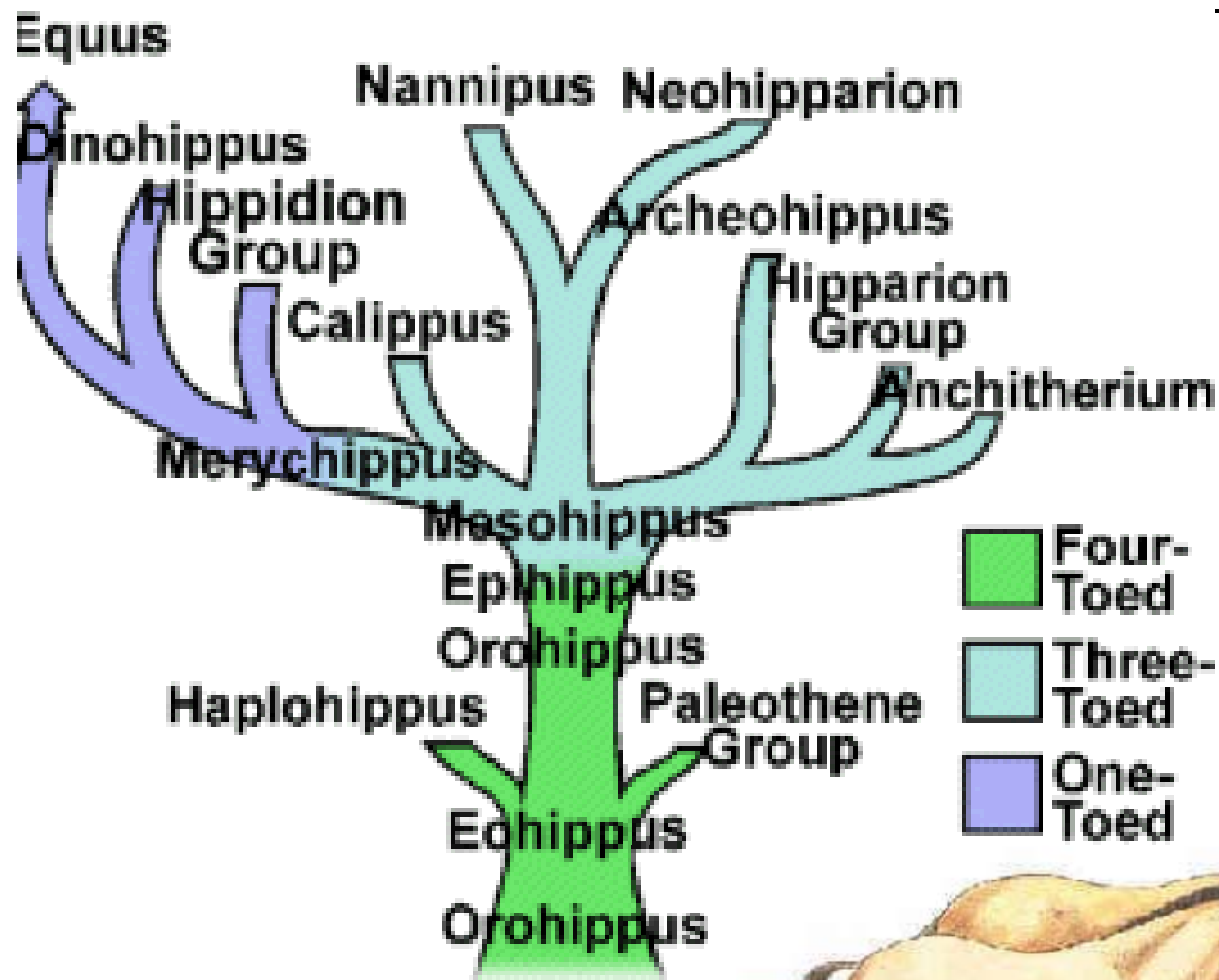


(b) The same adaptive radiation of *Anolis* has occurred on different islands, starting from different types of colonists.



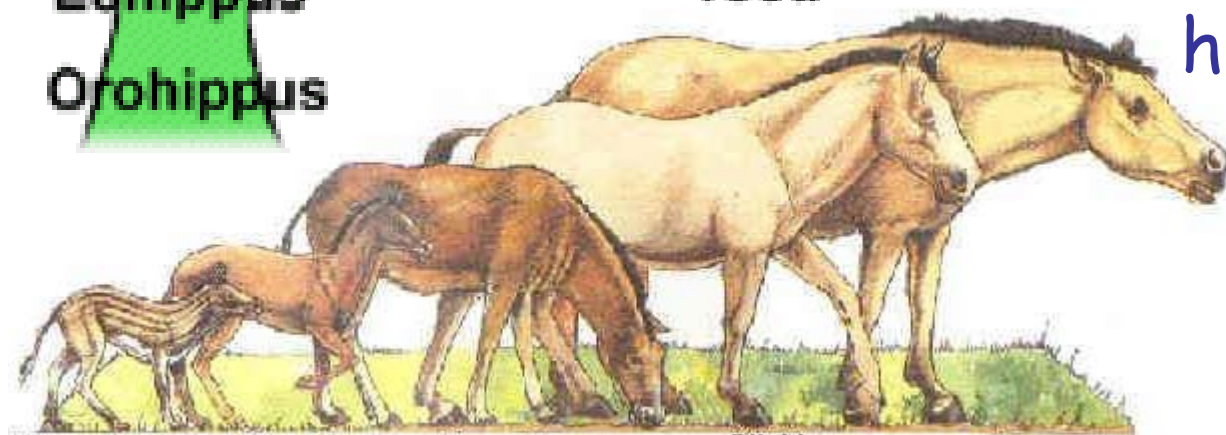
Copyright © 2008 Pearson Benjamin Cummings. All rights reserved.

Expand and diversify into areas without competition
(Area either 1) new or 2) former residents extinct.)



Most are
EXTINCT!

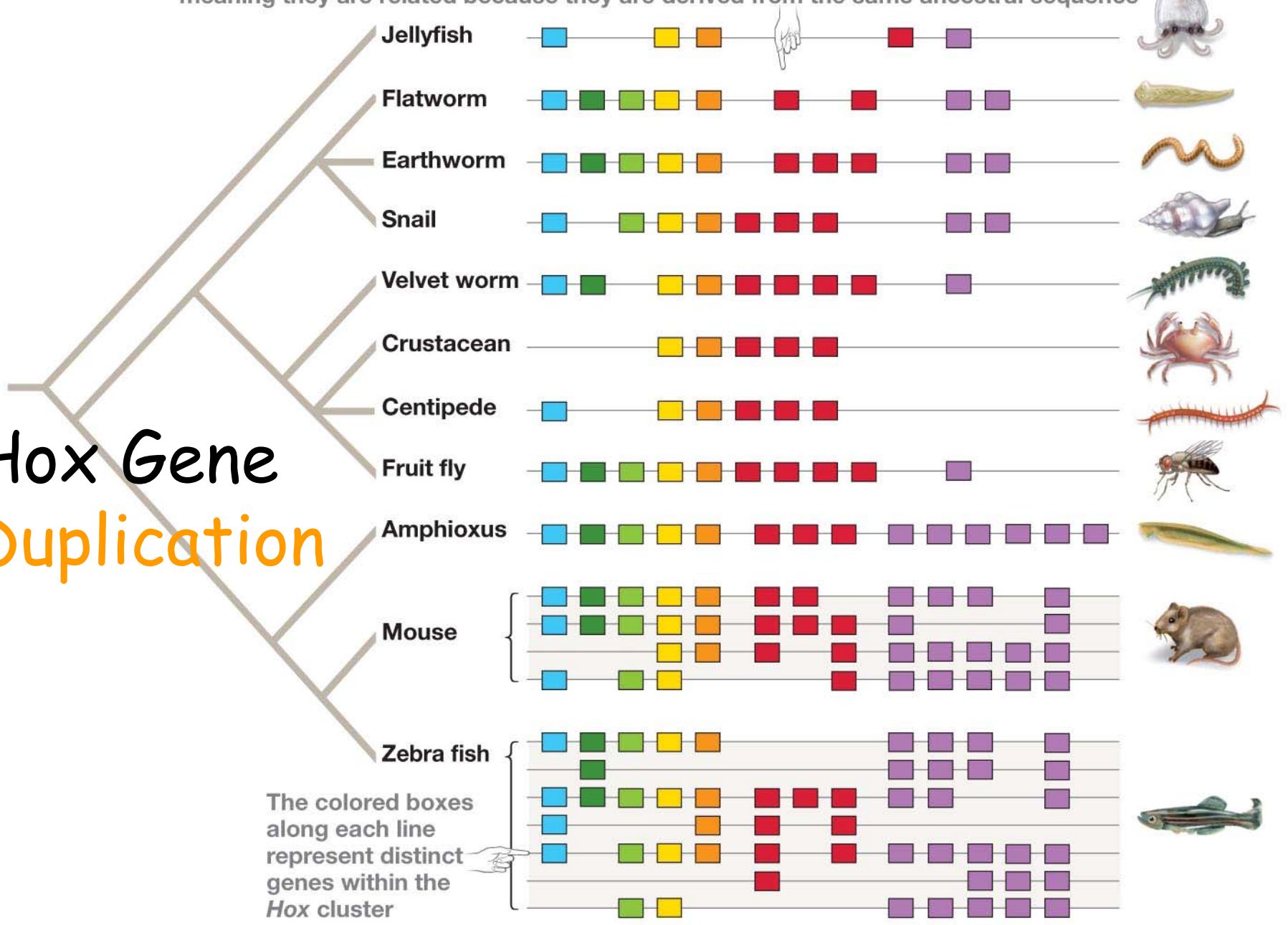
Modern
horse



Hyracotherium Mesohippus Merychippus Pliohippus Equus

The same-colored boxes within a vertical column are considered to be homologous—meaning they are related because they are derived from the same ancestral sequence

Hox Gene Duplication



The colored boxes along each line represent distinct genes within the *Hox* cluster

Living fossils: *Gingko*

(a) Triassic



(b)



Living fossils: Horseshoe crabs

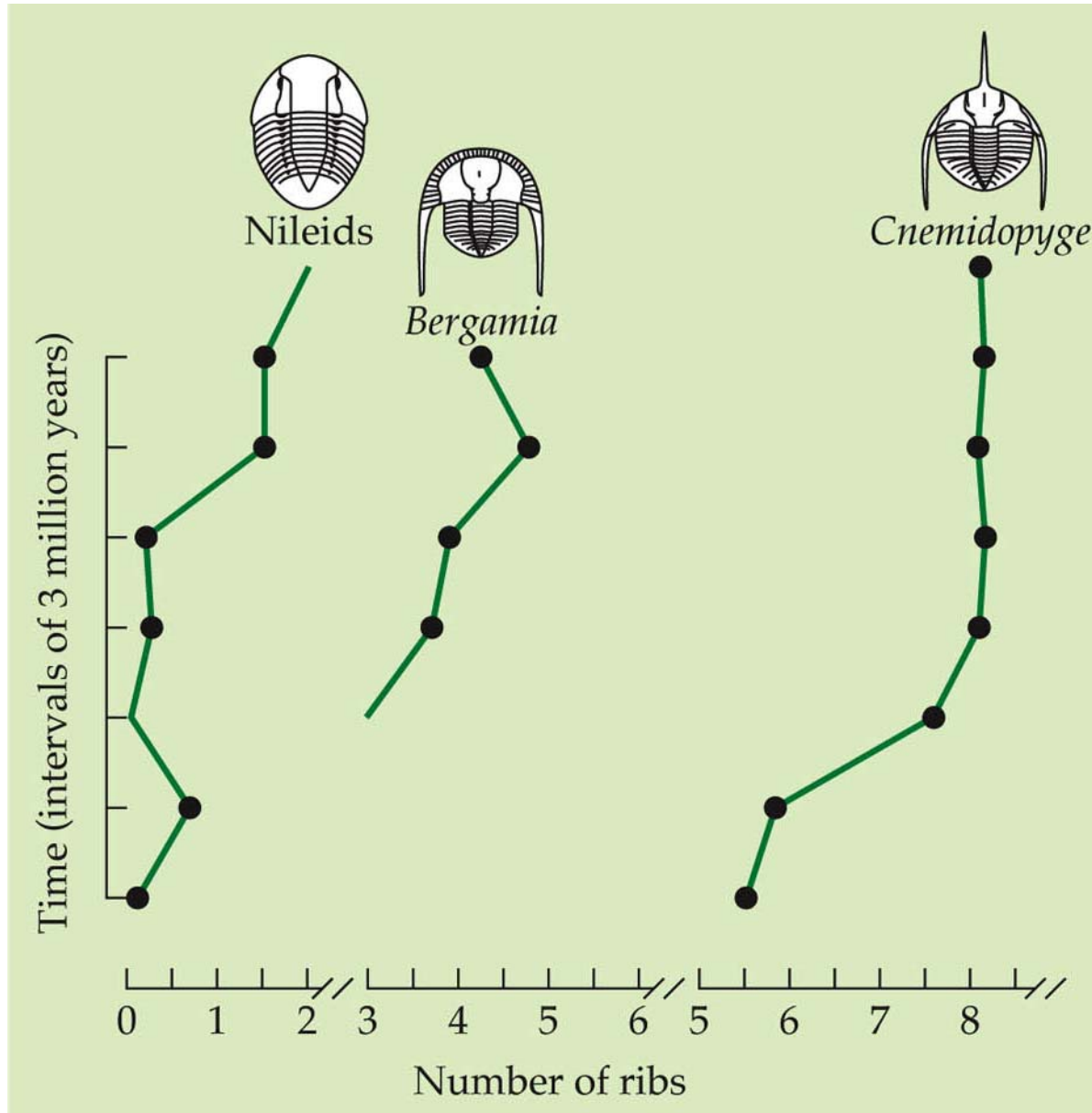


(b) *Limulus polyphemus*

~identical for
300 mya

sandy coastlines
are harsh
environments
that changed
little, so they
didn't need to
evolve anything
different

Gradual change: trilobite rib number



fossils show
number of ribs

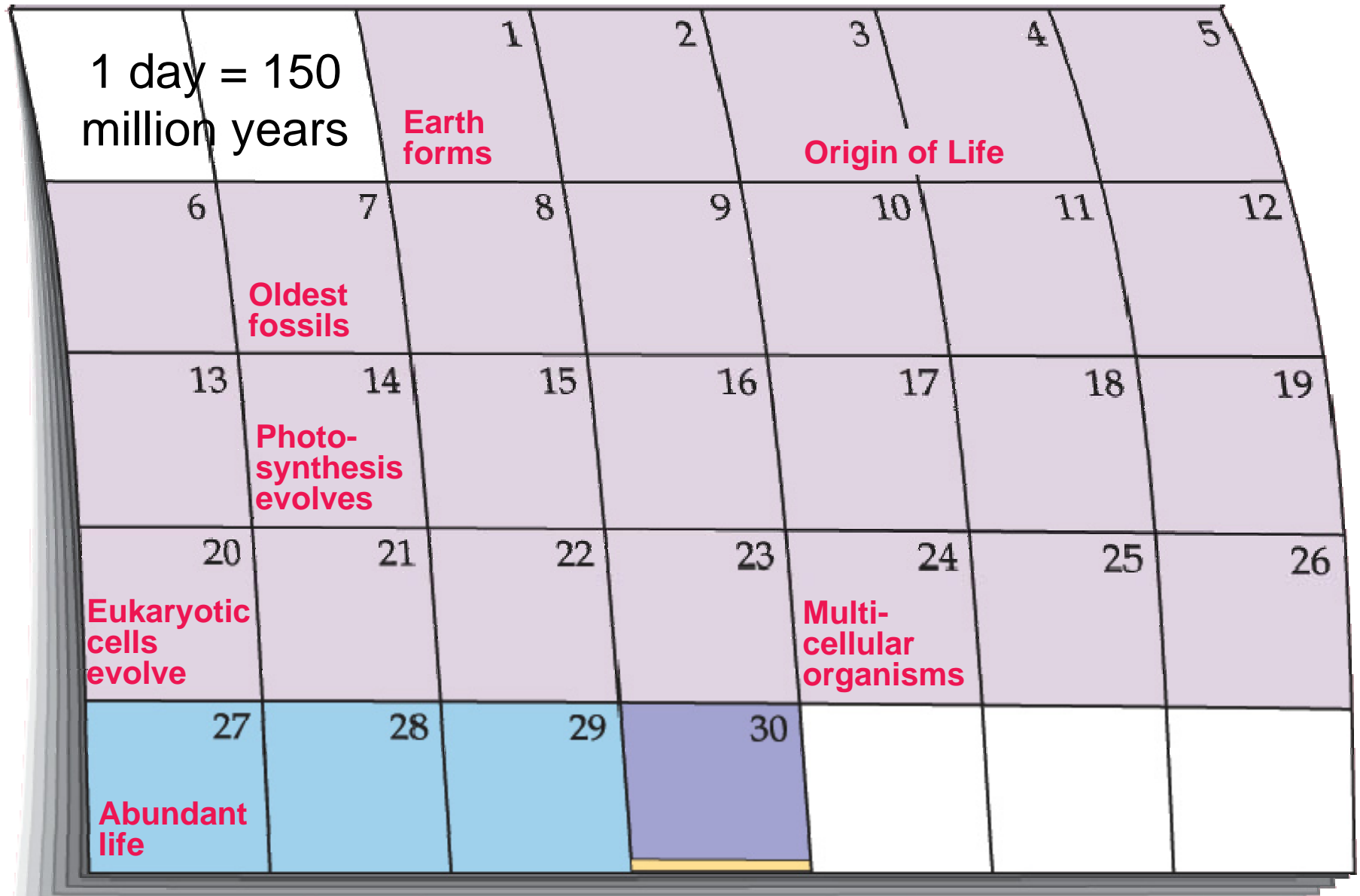
Biological history

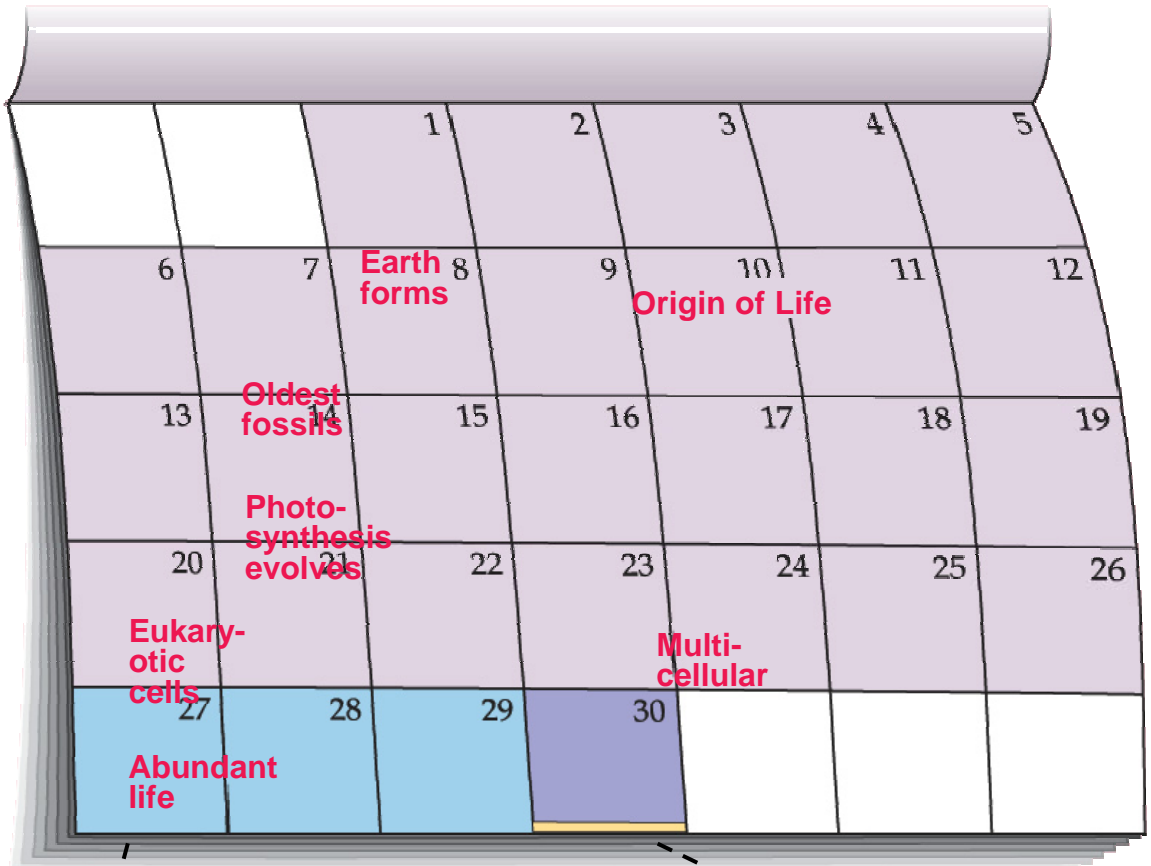
22.1 Earth's Geological History (Part 2)

RELATIVE TIME SPAN	ERA	PERIOD	ONSET	MAJOR EVENTS IN THE HISTORY OF LIFE
Precambrian	Cenozoic	Quaternary	1.8 mya ^a	Humans evolve; many large mammals become extinct
		Tertiary	65 mya	Diversification of birds, mammals, flowering plants, and insects
	Mesozoic	Cretaceous	144 mya	Dinosaurs continue to diversify; flowering plants and mammals diversify. Mass Extinction at end of period ($\approx 76\%$ of species disappear)
		Jurassic	206 mya	Diverse dinosaurs; radiation of ray-finned fishes
		Triassic	248 mya	Early dinosaurs; first mammals; marine invertebrates diversify; first flowering plants; Mass Extinction at end of period ($\approx 65\%$ of species disappear)
	Paleozoic	Permian	290 mya	Reptiles diversify; amphibians decline; Mass Extinction at end of period ($\approx 96\%$ of species disappear)
		Carboniferous	354 mya	Extensive "fern" forests; first reptiles; insects diversify
		Devonian	417 mya	Fishes diversify; first insects and amphibians. Mass Extinction at end of period ($\approx 75\%$ of species disappear)
		Silurian	443 mya	Jawless fishes diversify; first ray-finned fishes; plants and animals colonize land
		Ordovician	490 mya	Mass Extinction at end of period ($\approx 75\%$ of species disappear)
Precambrian	Cambrian	543 mya	Most animal phyla present; diverse algae	
	Precambrian		600 mya	Ediacaran fauna
			1.5 bya ^a	Eukaryotes evolve; several animal phyla appear
		3.8 bya	Origin of life; prokaryotes flourish	
		4.5 bya		

^amya, million years ago; bya, billion years ago.

Biological history





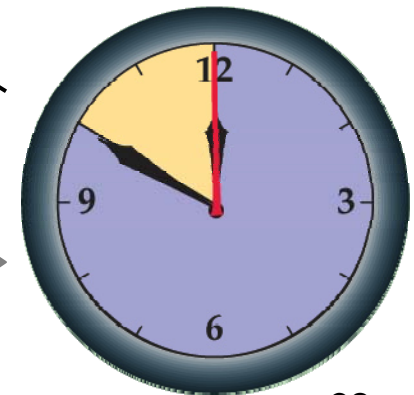
← **First hominids**
 ← *Homo sapiens*

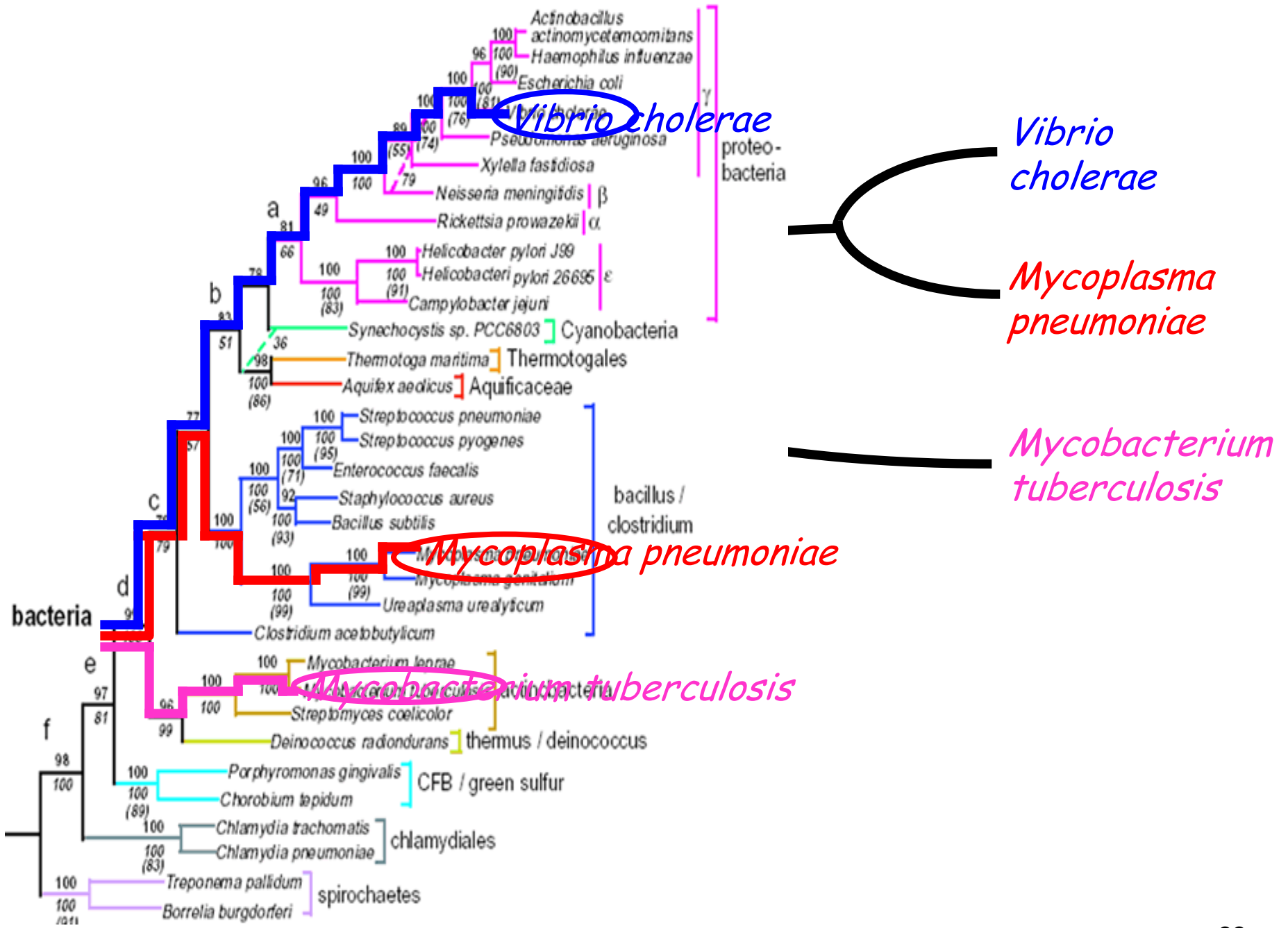
		1	2	3	4	5
		Earth forms		Origin of Life		
6	7	8	9	10	11	12
	Oldest fossils					
13	14	15	16	17	18	19
	Photo-synthesis evolves					
20	21	22	23	24	25	26
Eukary-otic cells			Multi-cellular			
27	28	29	30			
Abundant life						

Recorded history fills the last 5 seconds of day 30.

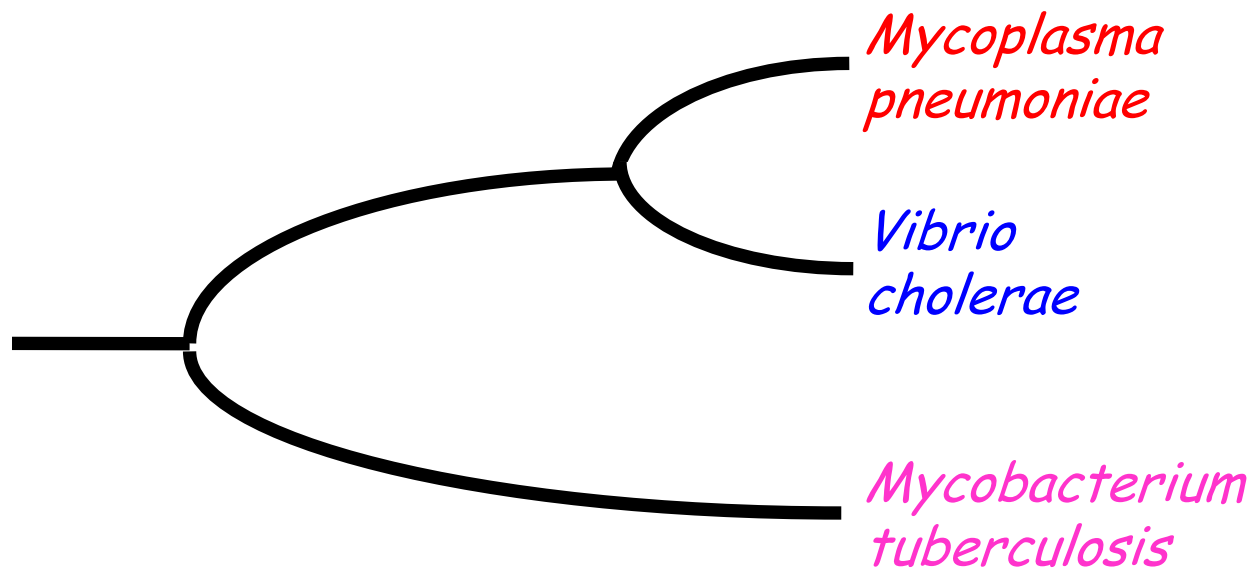
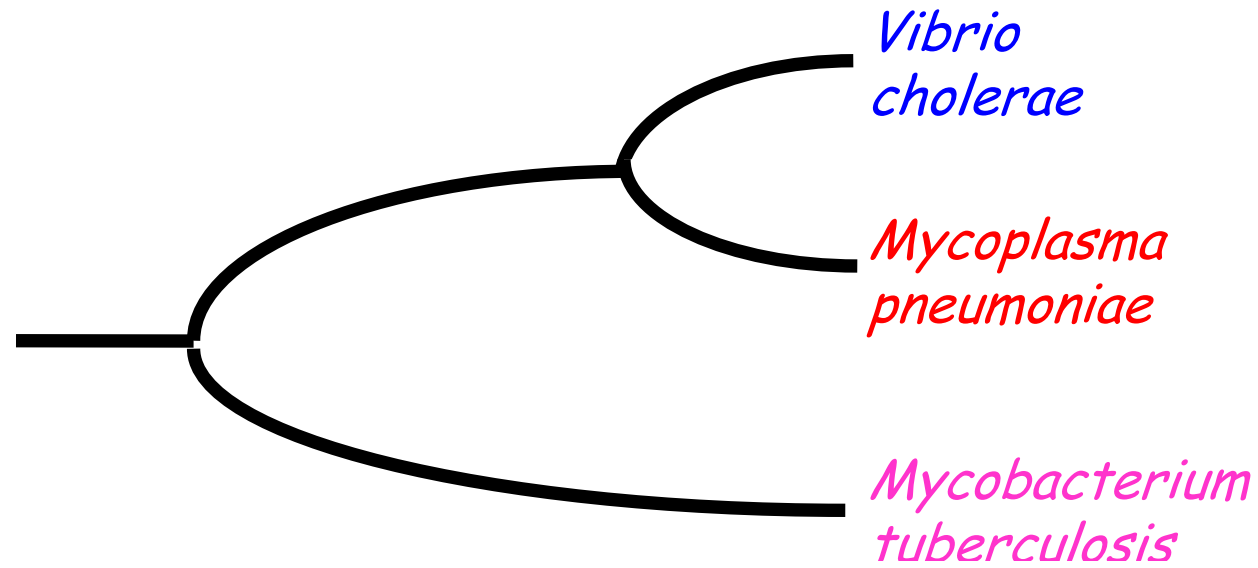
Modern humans appeared in the last 10 minutes of day 30.

27	28	29	30
Aquatic life	Land plants	Forests	Birds
Abundant fossils	Land animals	Insects	Flowering plants
		First mammals	Rise of Mammals
		Dinosaurs dominant	

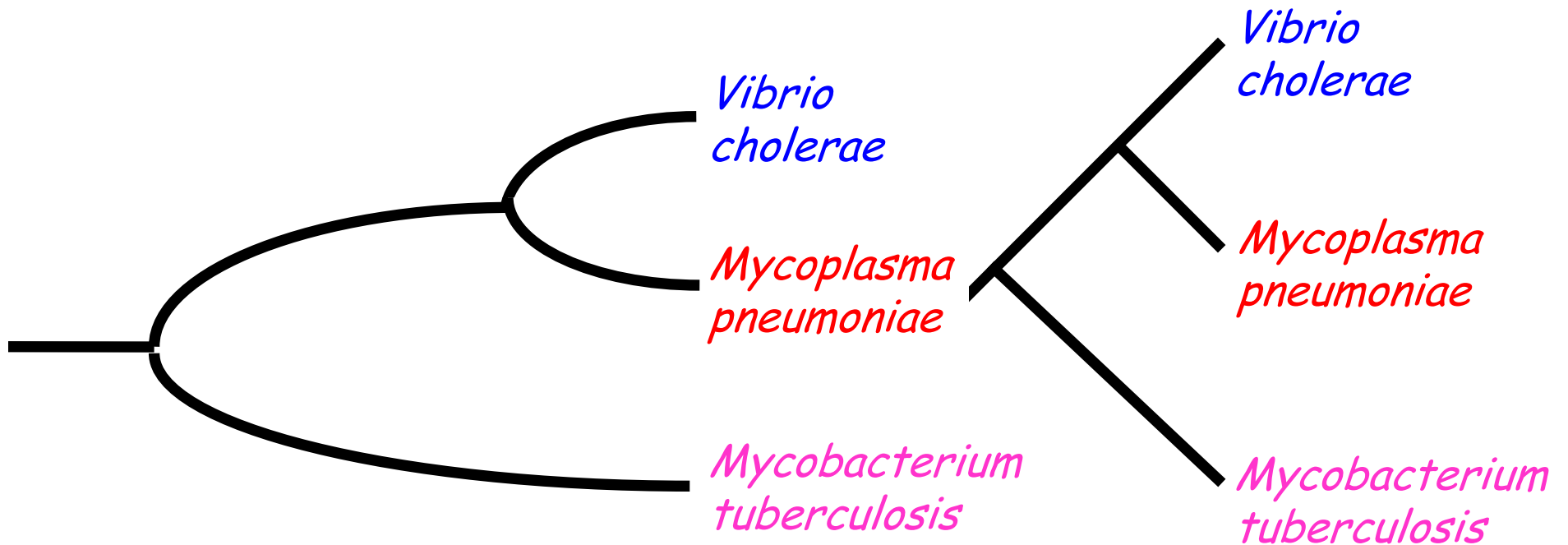




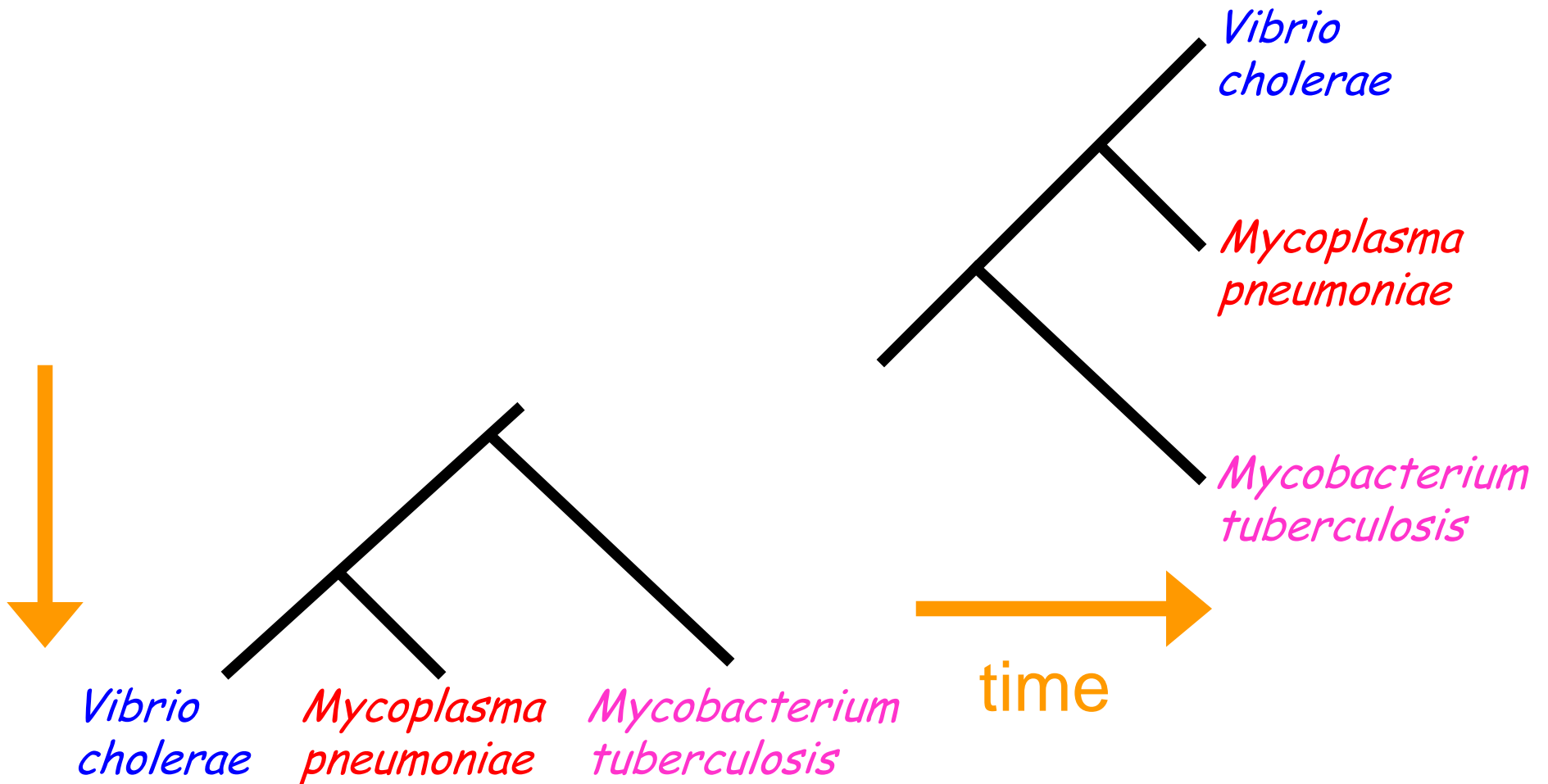
These trees are the same



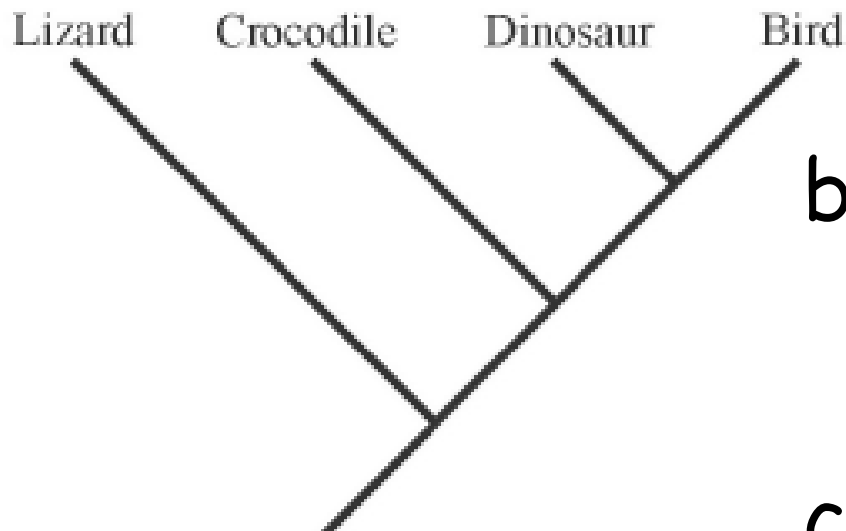
These trees are the same



These trees are the same

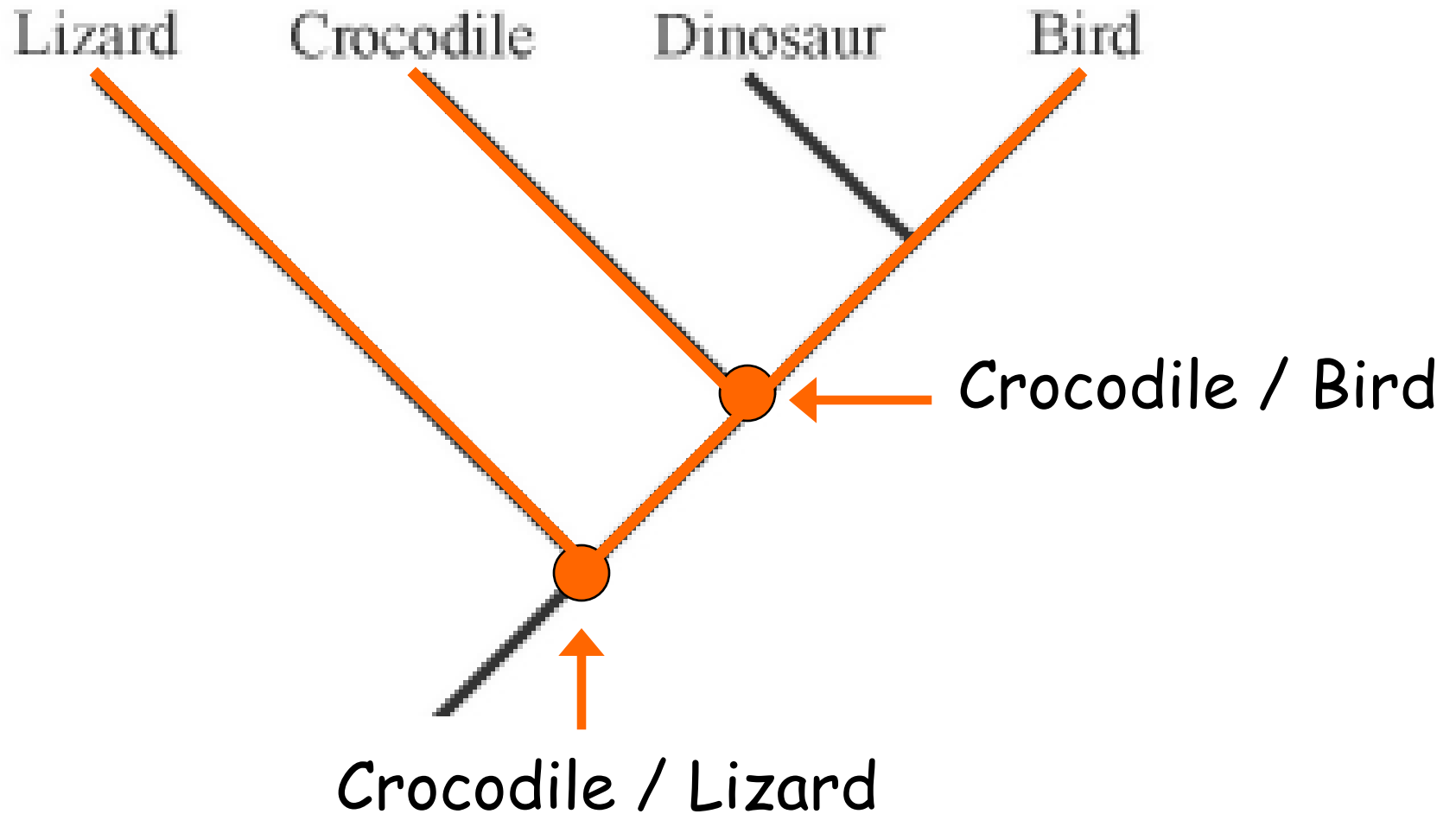


Which of these statements follows from the tree?

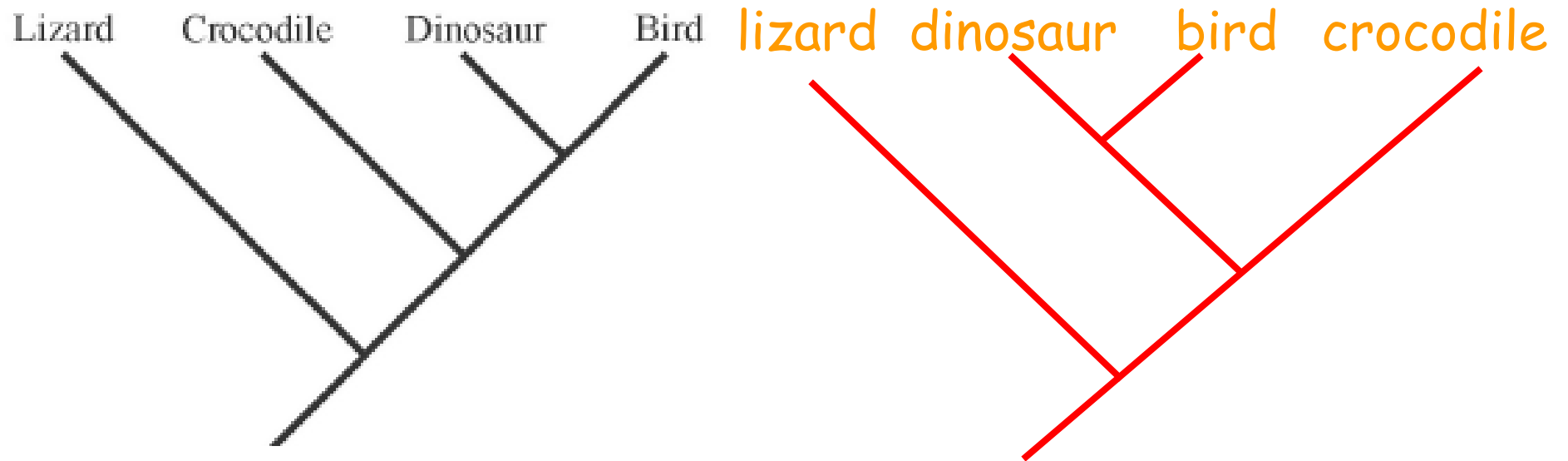


- a. Crocodiles are more closely related to lizards than to birds
- b. Crocodiles are more closely related to birds than to lizards
- c. Crocodiles are equally related to lizards and birds

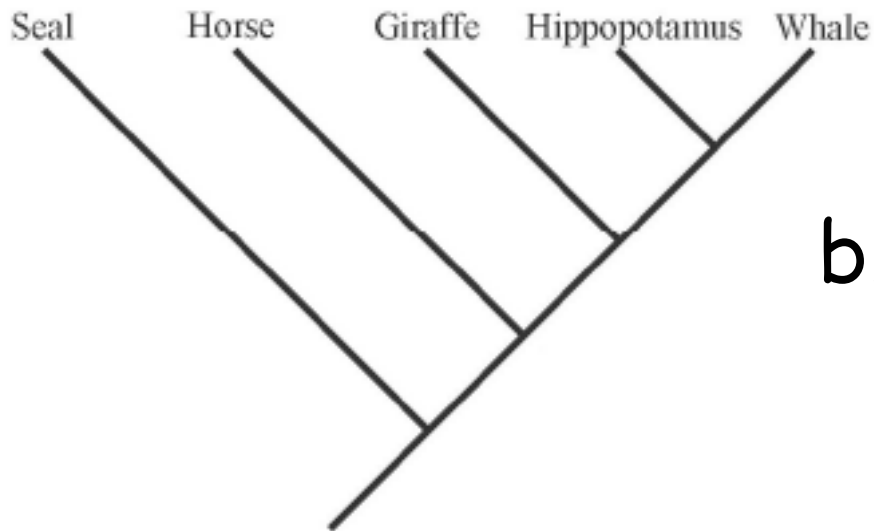
Focus on nodes = common ancestors



Another method: redraw tree with focus on crocodile



Which of these statements follows from the tree?



- a. A seal is more closely related to a horse than to a whale.
- b. A seal is more closely related to a whale than to a horse.
- c. A seal is equally related to a whale and a horse.

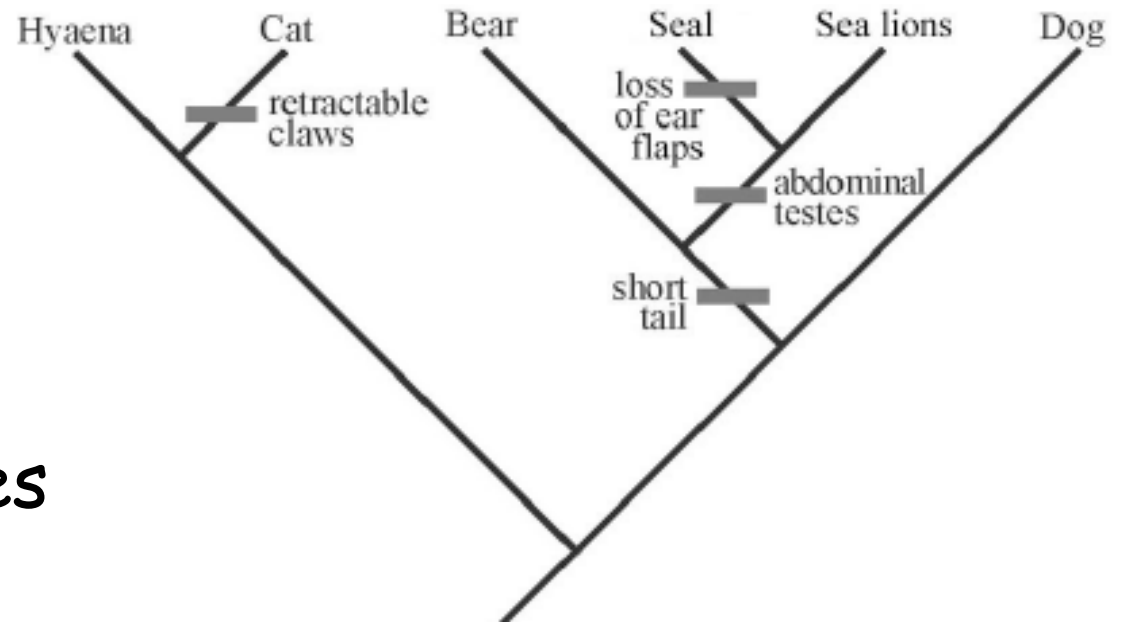
Ancestral traits

- Traits inherited from ancestor in distant past should be shared by large number of species
- Traits that first appeared in more recent ancestor should be shared by fewer species
- Ancestral trait = shared traits inherited from a common ancestor
- Derived trait = different from ancestral form

Nucleic acids are an ancestral trait

- Nucleic acids (DNA/RNA) as genetic material is ancestral to all life on Earth
- Specific genetic code is ancestral to most

The ancestor to this tree had a long tail, ear flaps, external testes, and fixed claws. Based on the tree, what traits does a **sea lion** have?



- a. long tail, ear flaps, external testes, and fixed claws
- b. short tail, no ear flaps, external testes, and fixed claws
- c. short tail, no ear flaps, abdominal testes, and fixed claws
- d. short tail, ear flaps, abdominal testes, and fixed claws
- e. long tail, ear flaps, abdominal testes, and retractable claws

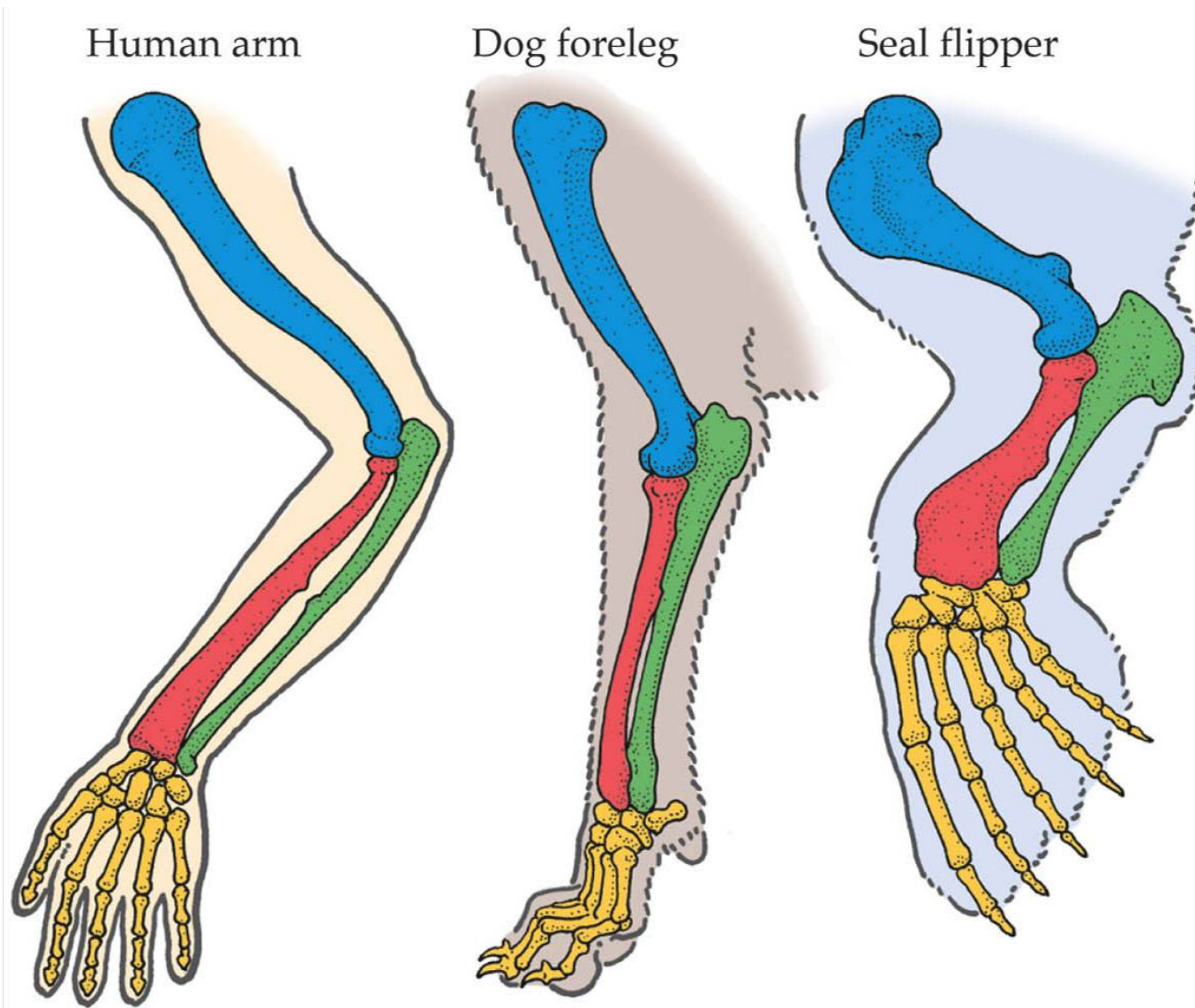
Ancestral vs derived may depend on scale of taxa

- In rodents: continuously growing incisors are **ancestral**: all rodents have them
- In mammals they are **derived**, unique to rodents

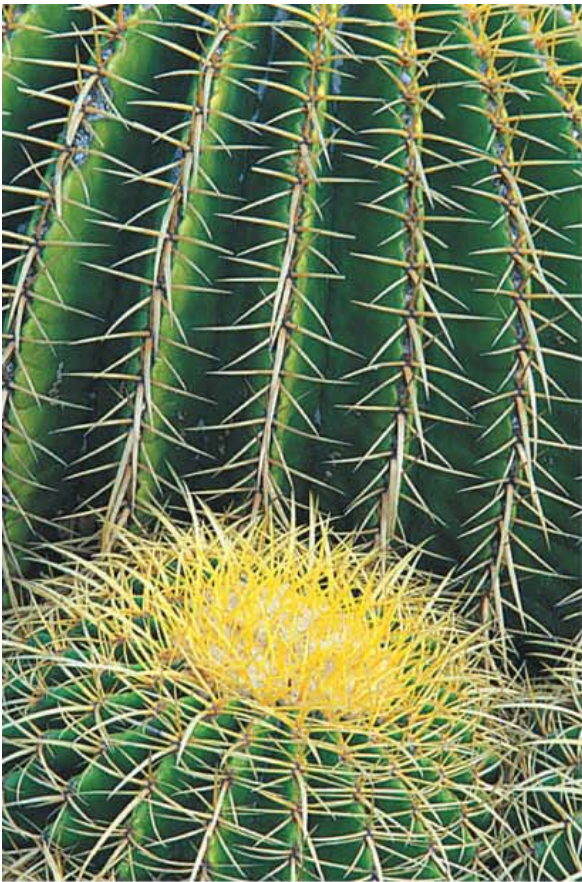
Homologous traits

- Features (DNA sequence, behavior, morphology) shared by species **descended from common ancestor** are called **homologous**
- e.g., vertebral column is homologous in vertebrates

Mammalian limbs are homologous, even with different functions



Homologous structures derived from leaves



Cheiridopsis tuberculata



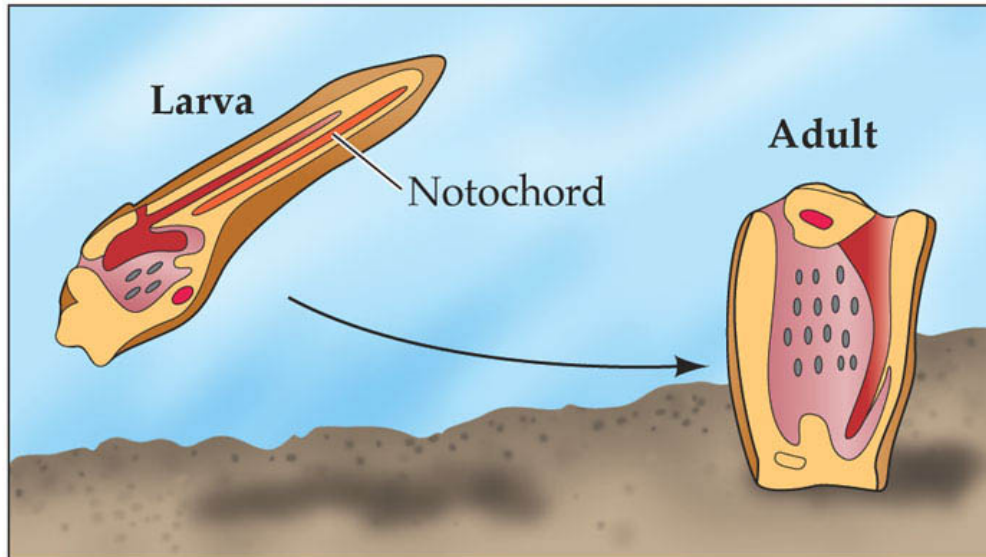
Heliconia sp.



Sarracenia purpurea

Some homologies are hidden by development

Sea squirt



Notochord homology seen only in early development, invisible in adult

Frog embryo



Both in Phylum Chordata

Deducing phylogeny

Look at shared traits

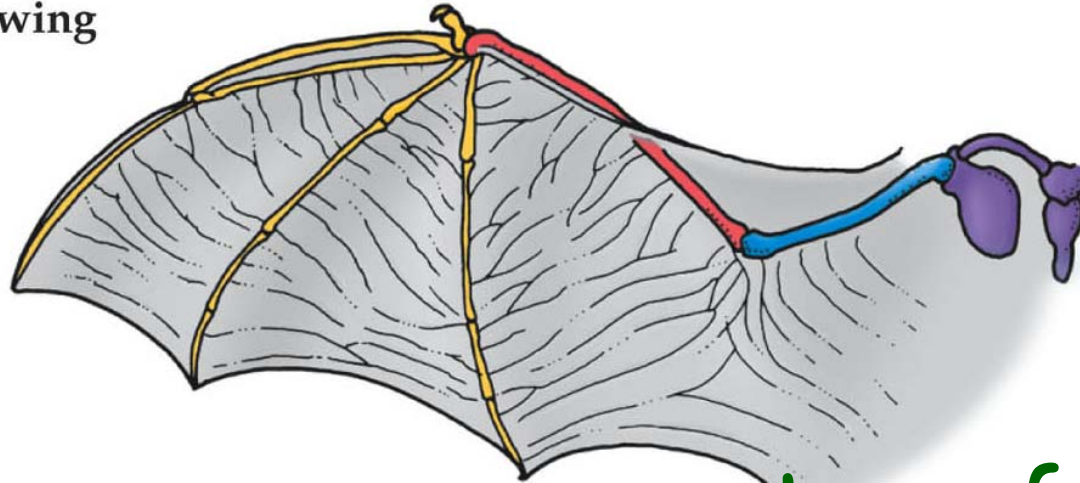
- What is ancestral state?
- How has it been **modified**?

Two processes make this difficult

- 1. **Convergent evolution** = similar selective pressures make **independently evolved traits look superficially similar**
- 2. **Evolutionary reversal** = character reverts from derived state back to ancestral state

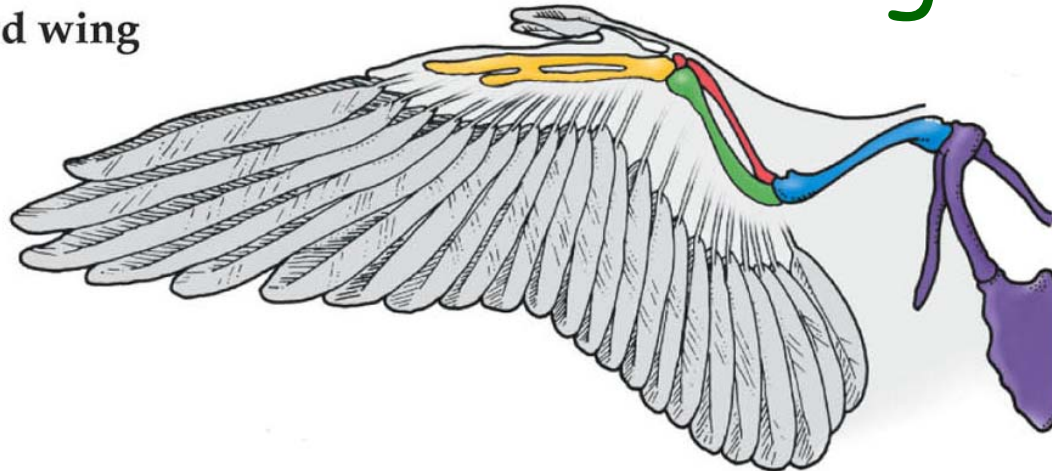
Bird and bat bones are homologous,
but not wings (convergent evolution)

Bat wing



Did the common ancestor of birds
and bats have wings?

Bird wing



Homoplastic traits

Convergent evolution and evolutionary reversal generate **homoplasies**: traits that are similar for some reason other than inheritance from a common ancestor

Bird and bat wings are **homoplastic**

Phrynosoma asio,
Long-spined
Horned Lizard
MEXICO



© Wendy Hodges

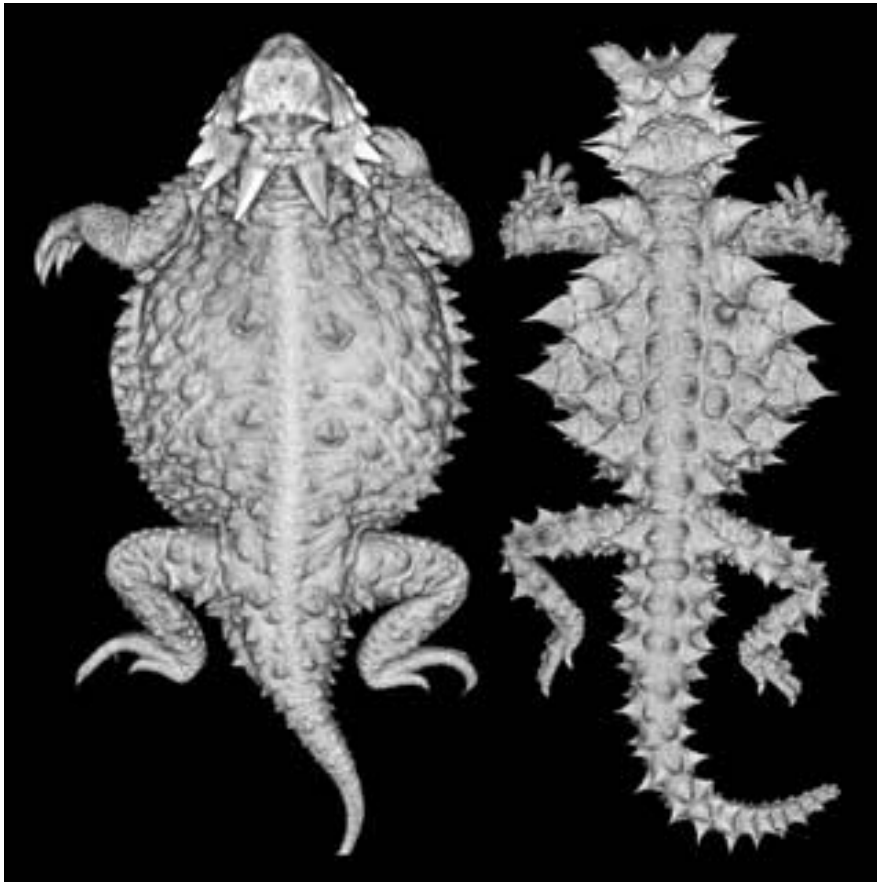


http://digimorph.org/specimens/Phrynosoma_asio/

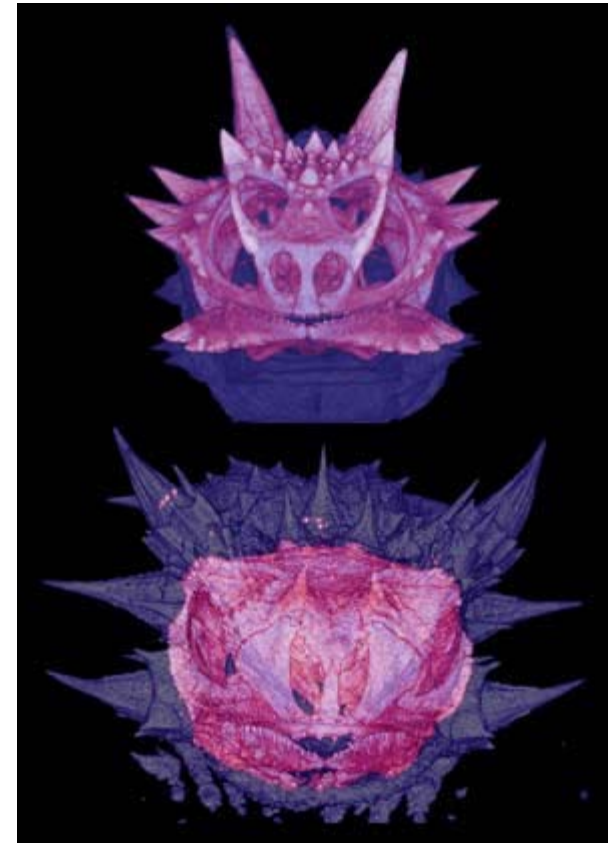
Moloch horridus, Thorny Devil, AUSTRALIA



Spiny head: **Homoplasy** or Homology?



Phrynosoma cornutum (left) vs. *Moloch horridus* (right)



P. cornutum (top) vs. *M. horridus* (bottom);
semi-transparent flesh in blue, bone in red

All of the following can result in homoplasy except

- a. similar selection pressures.
- b. reverse evolution.
- c. parallel evolution.
- d. descent from a recent common ancestor.
- e. convergent evolution.

Creating a phylogeny









1. Choose the characters and identify the possible forms (traits) of the characters
2. Determine ancestral and derived traits
3. Distinguish homologous from homoplastic traits









Traditional character choice

- Most often **morphology**
 - describes size and shape of body parts
 - can be seen directly in fossil record
- Also physiological, behavioral, molecular and structural traits as available
- The more characters measured, the more the inferred phylogenies should converge on each other and on the real evolutionary pattern

Simple phylogeny example









- Choose 8 vertebrate species and hypothesize their phylogeny
- Traits either present (+) or absent (-)
- Assume that each derived trait evolved only once and that no derived traits were lost

TAXON	JAWS	LUNGS	CLAWS OR NAILS	GIZZARD
 Lamprey	—	—	—	—
 Perch	+	—	—	—
 Salamander	+	+	—	—
 Lizard	+	+	+	—
 Crocodile	+	+	+	+
 Pigeon	+	+	+	+
 Mouse	+	+	+	—
 Chimpanzee	+	+	+	—

TAXON	FEATHERS	FUR	MAMMARY GLANDS	KERATINOUS SCALES
 Lamprey	—	—	—	—
 Perch	—	—	—	—
 Salamander	—	—	—	—
 Lizard	—	—	—	+
 Crocodile	—	—	—	+
 Pigeon	+	—	—	+
 Mouse	—	+	+	—
 Chimpanzee	—	+	+	—

Determine ancestral and derived traits

- Chimp and mouse share mammary glands and fur, other animals lack them
- Ancestral or derived? **derived**

TAXON	FEATHERS	FUR	MAMMARY GLANDS	KERATINOUS SCALES
 Lamprey	-	-	-	-
 Perch	-	-	-	-
 Salamander	-	-	-	-
 Lizard	-	-	-	+
 Crocodile	-	-	-	+
 Pigeon	+	-	-	+
 Mouse	-	+	+	-
 Chimpanzee	-	+	+	-

Determine ancestral and derived traits

- Use similar reasoning on other traits
- Lamprey has no derived traits, so it is an **outgroup**

Your Turn...

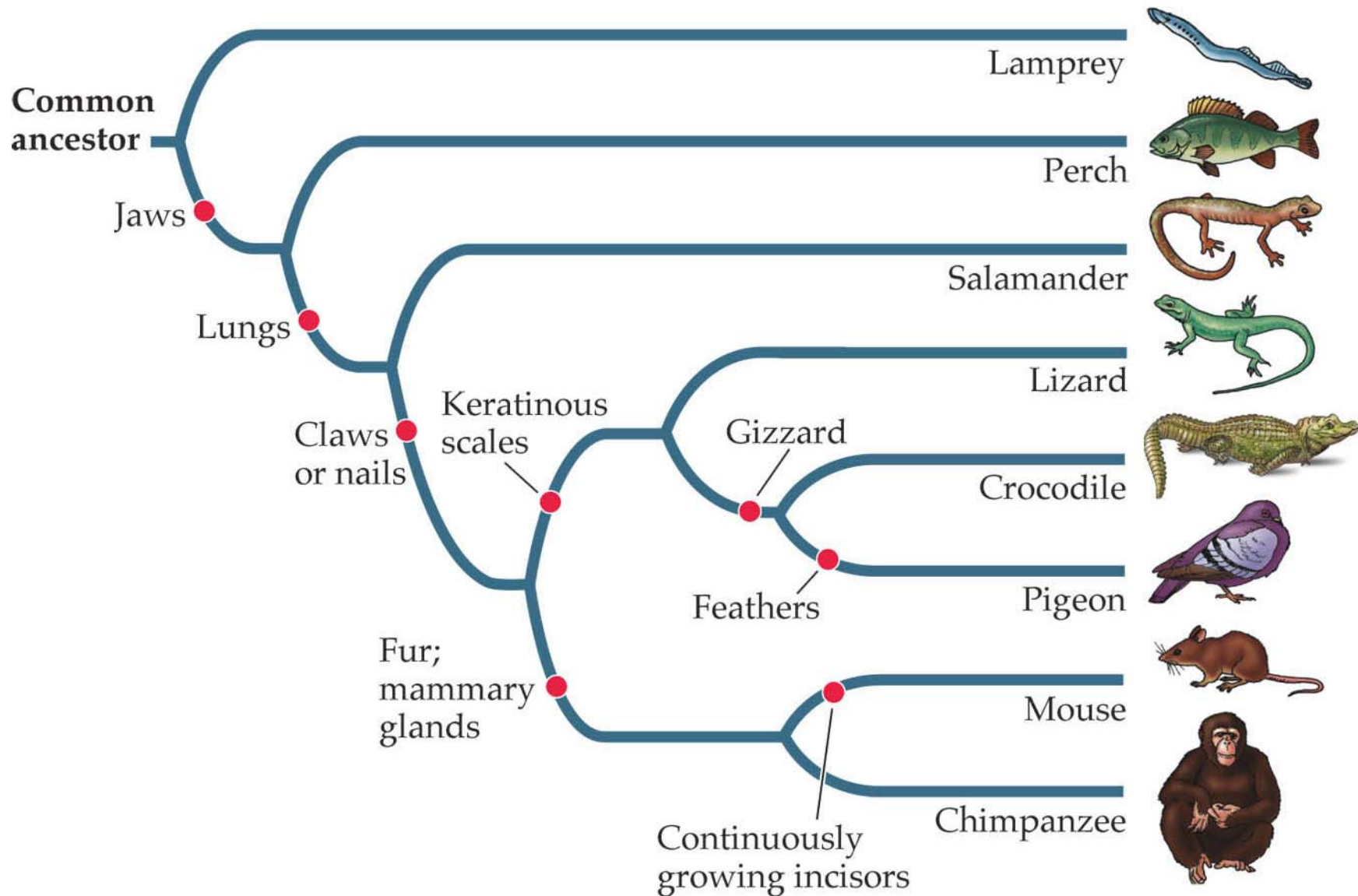


TAXON	DERIVED TRAIT ^a			
	JAWS	LUNGS	CLAWS OR NAILS	GIZZARD
Lamprey	-	-	-	-
Perch	+	-	-	-
Salamander	+	+	-	-
Lizard	+	+	+	-
Crocodile	+	+	+	+
Pigeon	+	+	+	+
Mouse	+	+	+	-
Chimpanzee	+	+	+	-



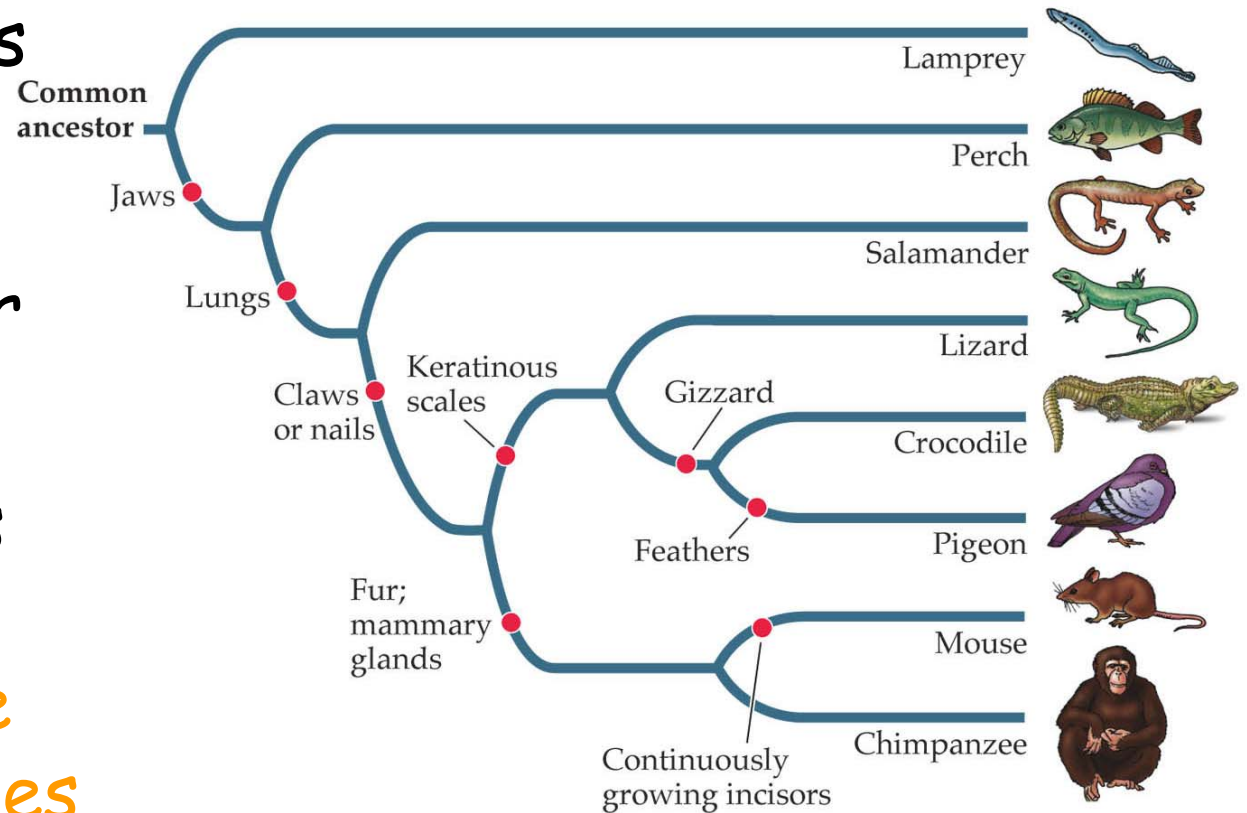
TAXON	FEATHERS	FUR	MAMMARY GLANDS	KERATINOUS SCALES
Lamprey	-	-	-	-
Perch	-	-	-	-
Salamander	-	-	-	-
Lizard	-	-	-	+
Crocodile	-	-	-	+
Pigeon	+	-	-	+
Mouse	-	+	+	-
Chimpanzee	-	+	+	-

Simple phylogeny example



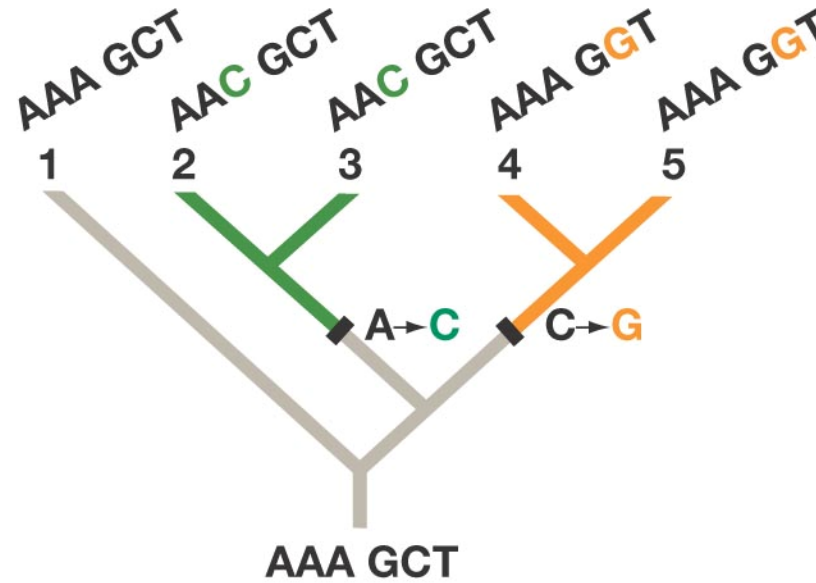
Not always so simple

- Assumed derived traits appeared once and were never lost
- If we had included snakes, this wouldn't have worked: they have scales but no claws
- Snakes lost limbs and claws during descent from a common ancestor with lizards
- Parsimony means we assume the smallest possible number of changes

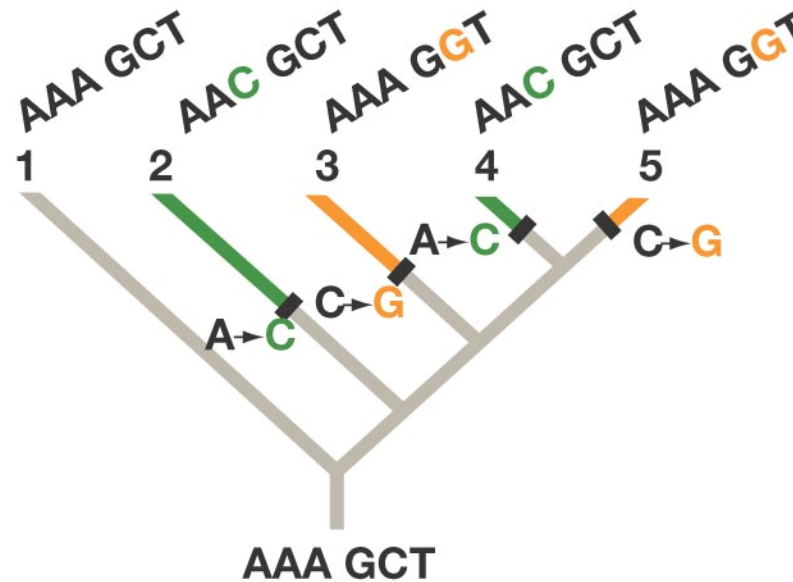


Parsimony

(a) Two changes



(b) Four changes



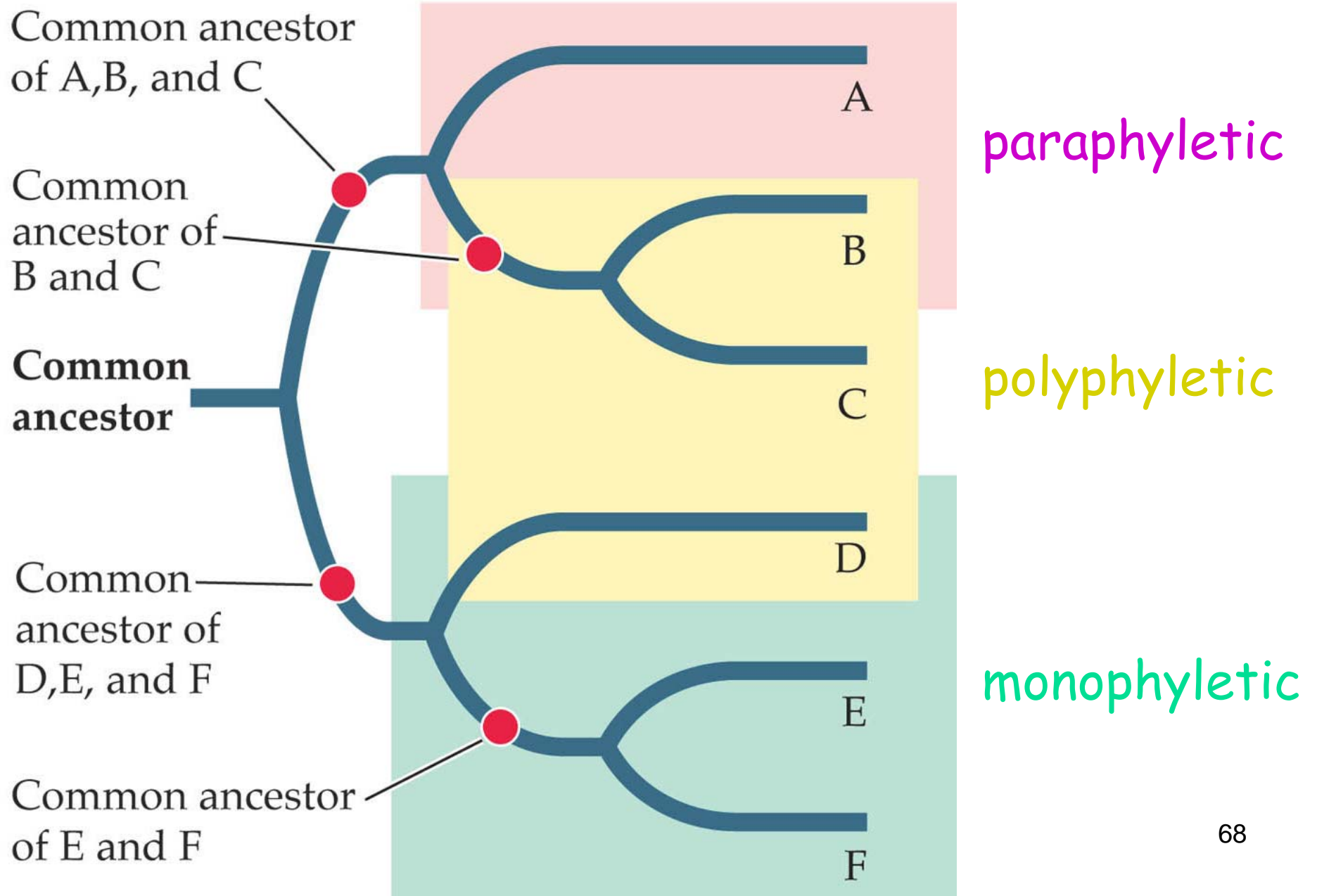
Modern character choice

Today, most phylogenies come from
gene sequences

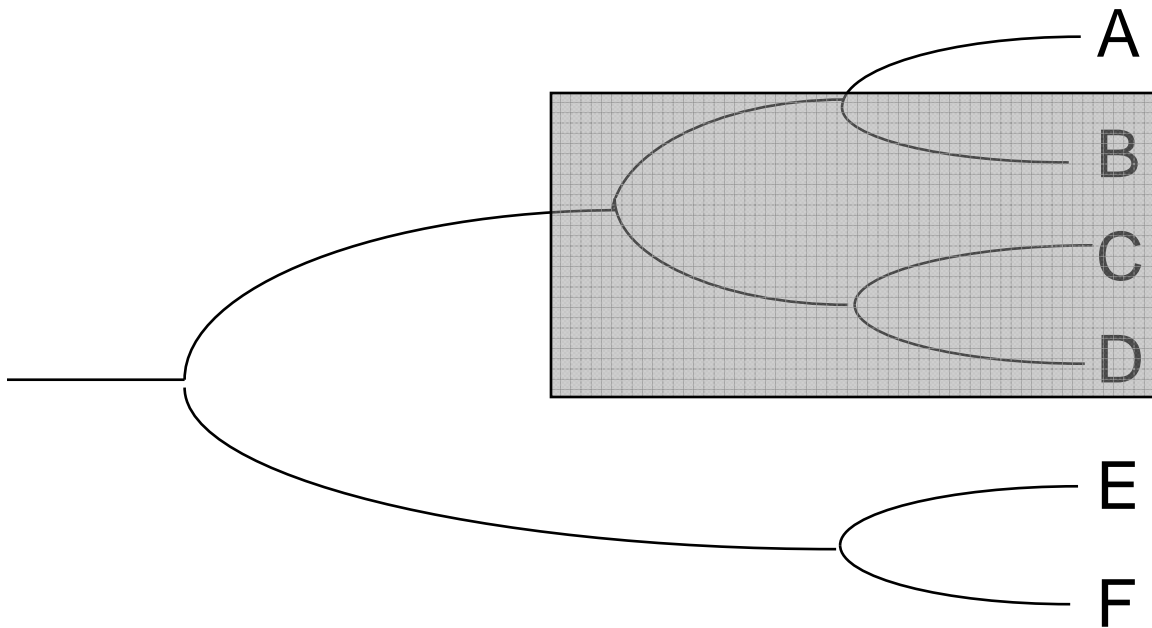
- DNA/RNA/protein

Classification should reflect evolutionary relationships

- A **monophyletic** group (or **clade**) **best** contains all descendents of a particular ancestor and no other organisms
- A **polyphyletic** taxon contains members with more than one recent common ancestor
- A **paraphyletic** group contains some, but not all of the descendents of a particular ancestor

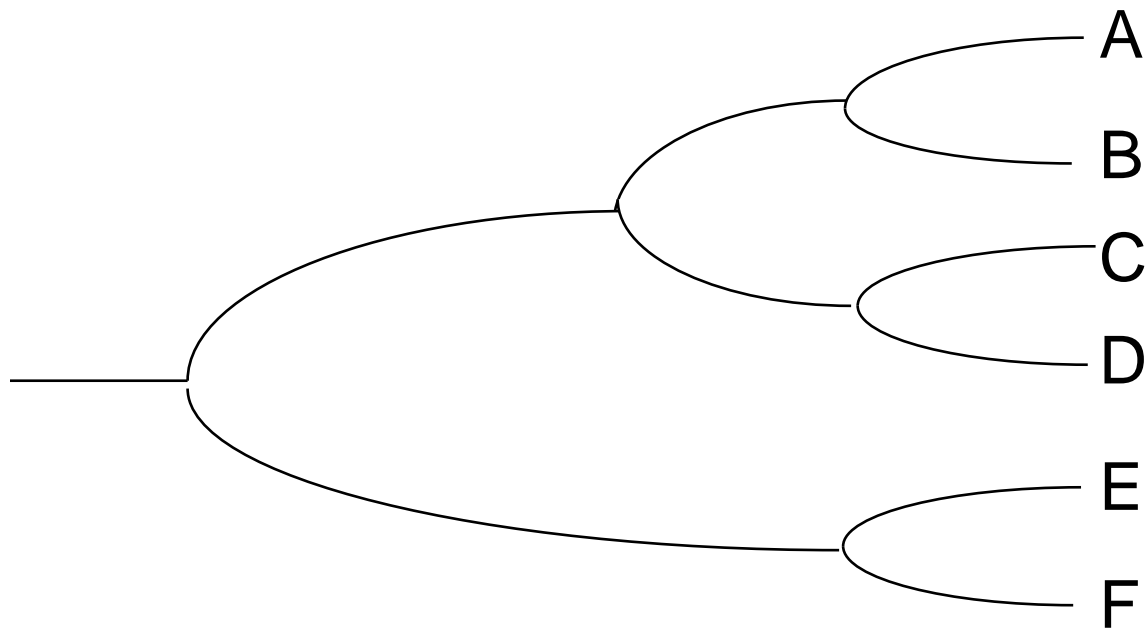


A group consisting of the shaded species is best described as



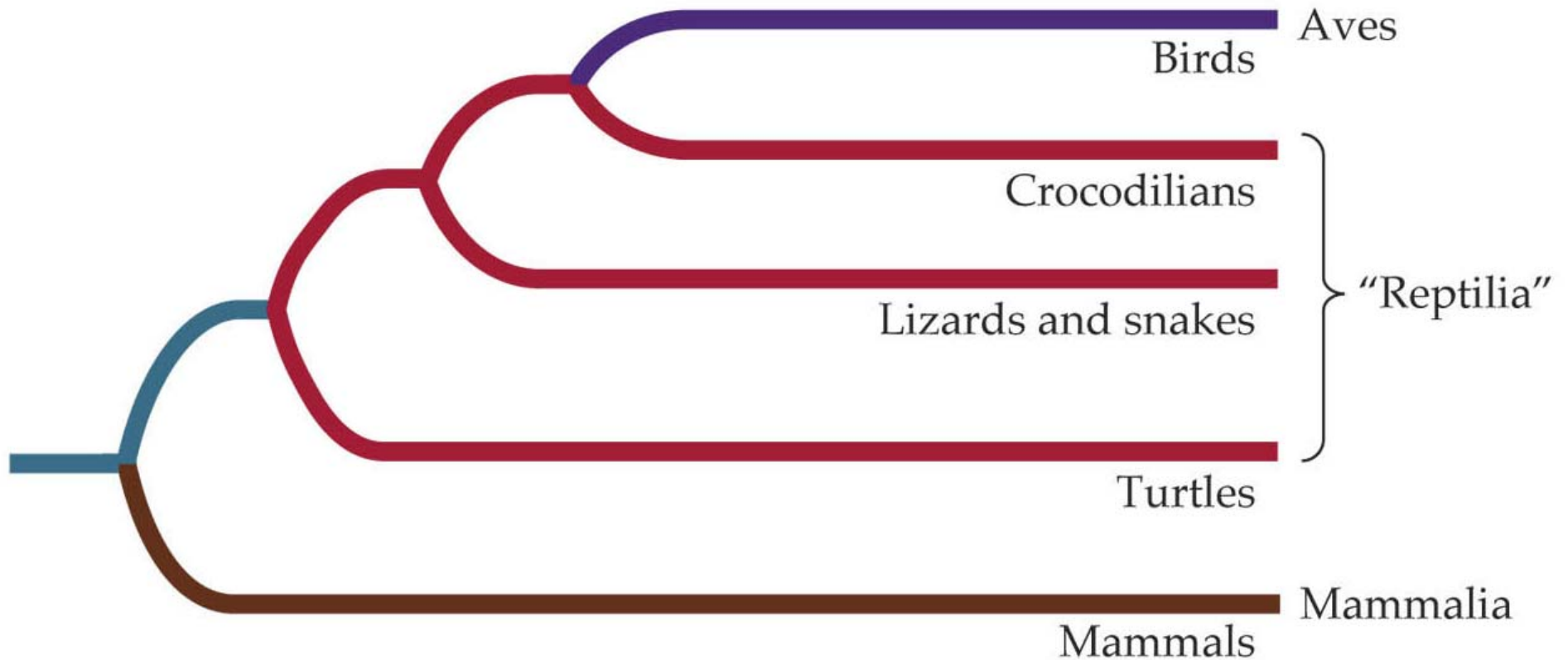
- a. polyphyletic.
- b. paraphyletic.
- c. monophyletic.

A particular trait is found only in species A and E.
Assuming that the phylogenetic tree is correct,
the simplest model describes this trait as



- a. orthologous.
- b. ancestral.
- c. homoplastic.
- d. homologous.

Should reptiles be a group?



Molecular evidence shows birds and crocodilians closer than crocodilians and other reptiles

Reptiles are **paraphyletic**, since birds excluded

Birds look very different

- Birds have rapidly evolved unique derived traits since they separated from reptiles
- Groups called **grades** have changed rapidly. May be an appropriate group even if paraphyletic
- General tendency to eliminate paraphyletic groups as we learn more, but some familiar categories, such as reptiles, won't disappear in a hurry

Uses of phylogenetic trees

1. How are different species related?
2. What traits do you expect a newly discovered species to have?
3. How many times has a trait evolved?
4. Which molecular change is responsible for adaptation?
5. When did lineages split?
use DNA sequence and the "molecular clock"

The Evolution of Macromolecules

- The **neutral theory** of molecular evolution:
 - postulates that, at the molecular level, the **majority of mutations are selectively neutral.**
- Thus, macromolecule evolution, and much of the genetic variation within species, does not result from positive selection of advantageous alleles nor stabilizing selection.
- Mutation fixation rate is theoretically constant and equal to the neutral mutation rate - a molecular clock.
 - The concept of the **molecular clock** states that **macromolecules should diverge from one another at a constant rate.**

Biological Clock

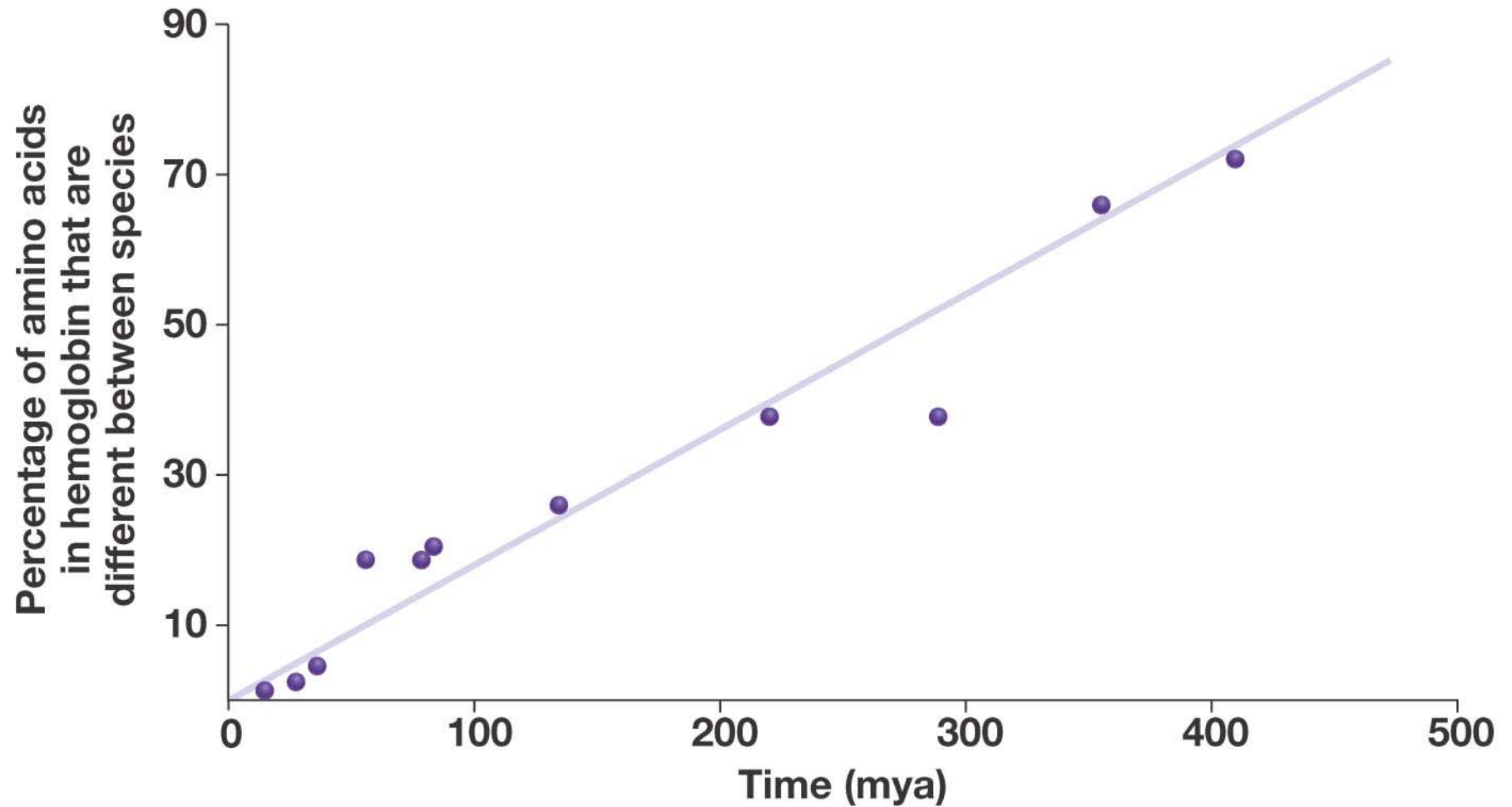
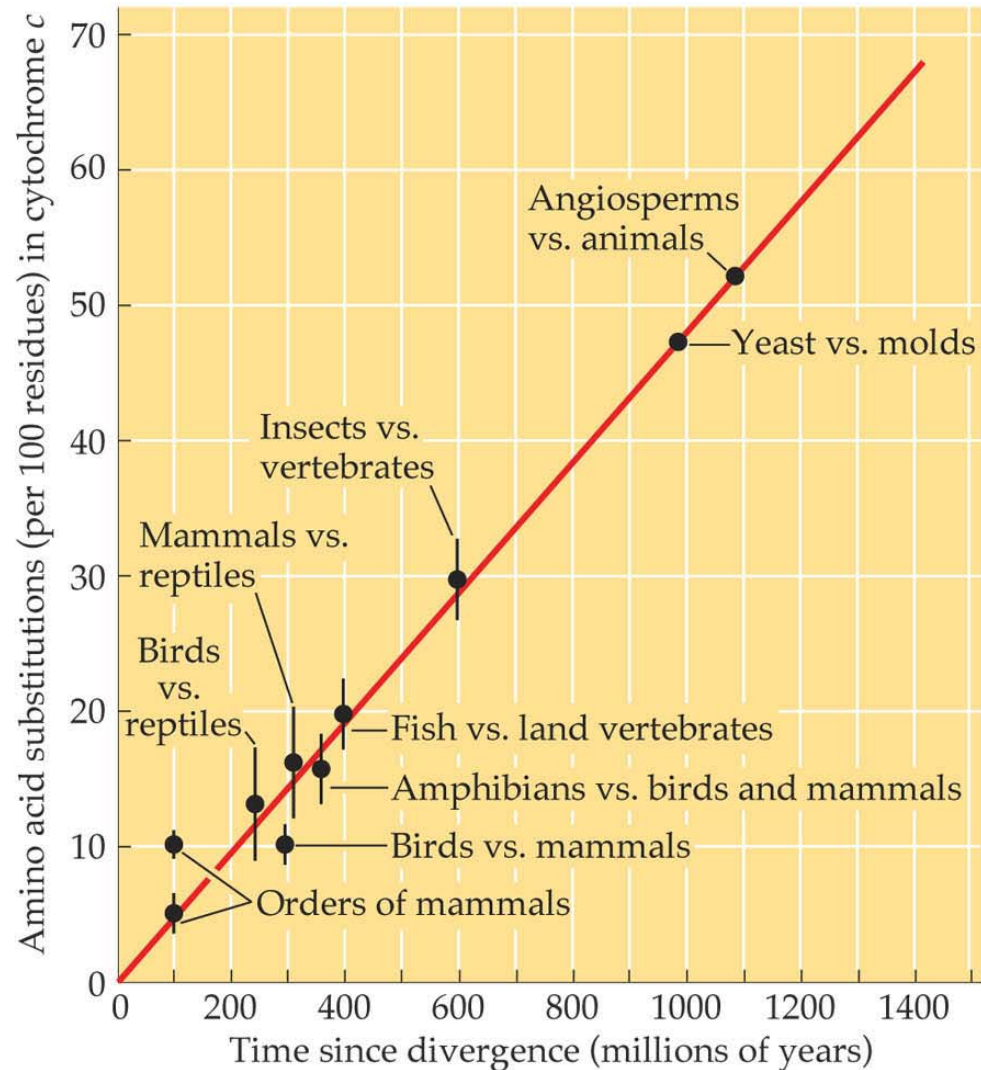


Figure 26.5 *Cytochrome c Has Evolved at a Constant Rate*

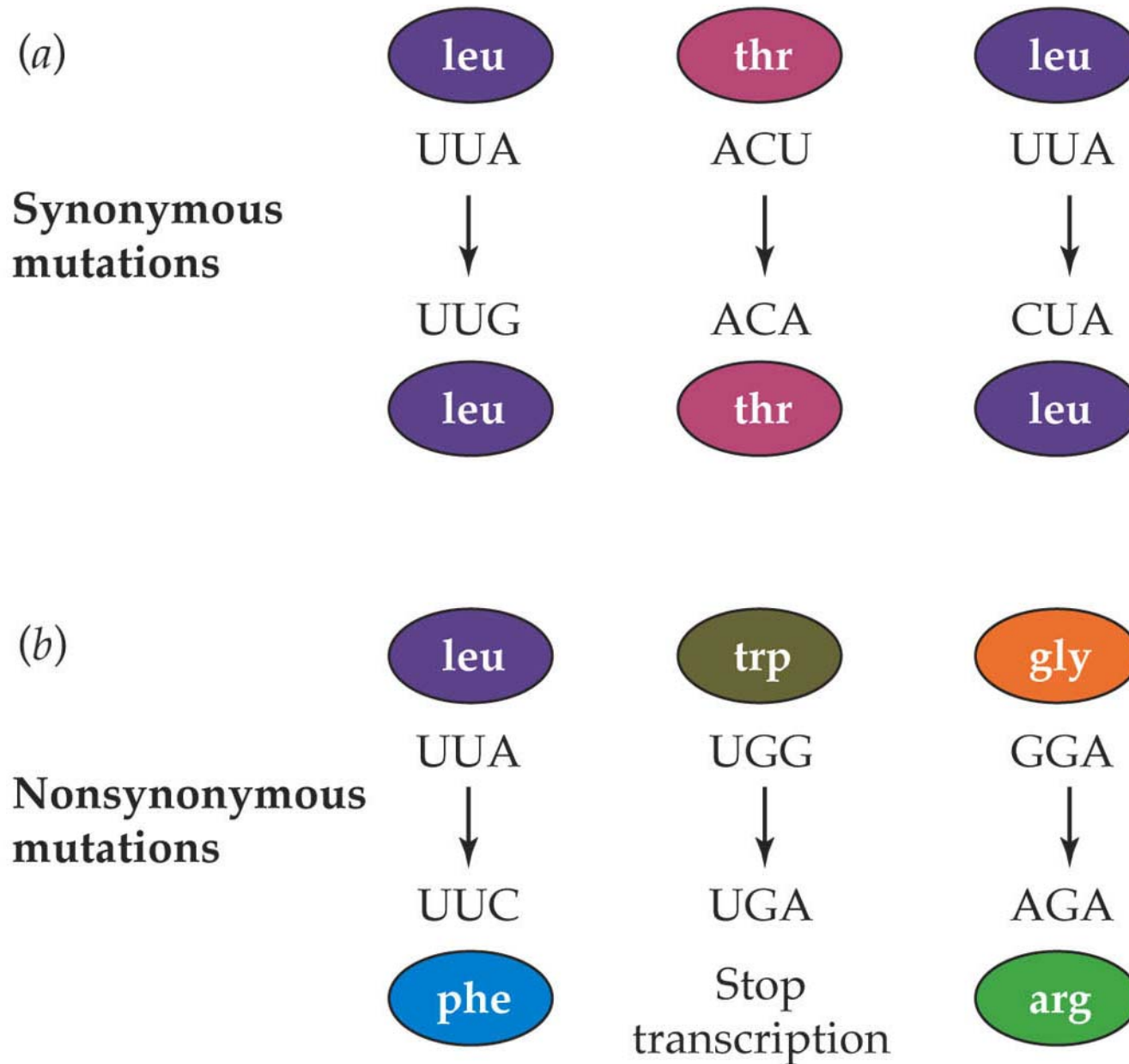


Cytochrome *c* sequences have evolved at a relatively **constant rate**.

The Evolution of Macromolecules

- Molecular evolution differs from phenotypic evolution in one important way:
 - In addition to **natural selection**, **random genetic drift** and **mutation** exert important influences on the rates and directions of molecular evolution.
 - What does this say about our study of "Evolutionary Theory"?
- Many mutations, called **silent or synonymous** mutations, do not alter the proteins they encode
- A **nonsynonymous** mutation does change the amino acid sequence

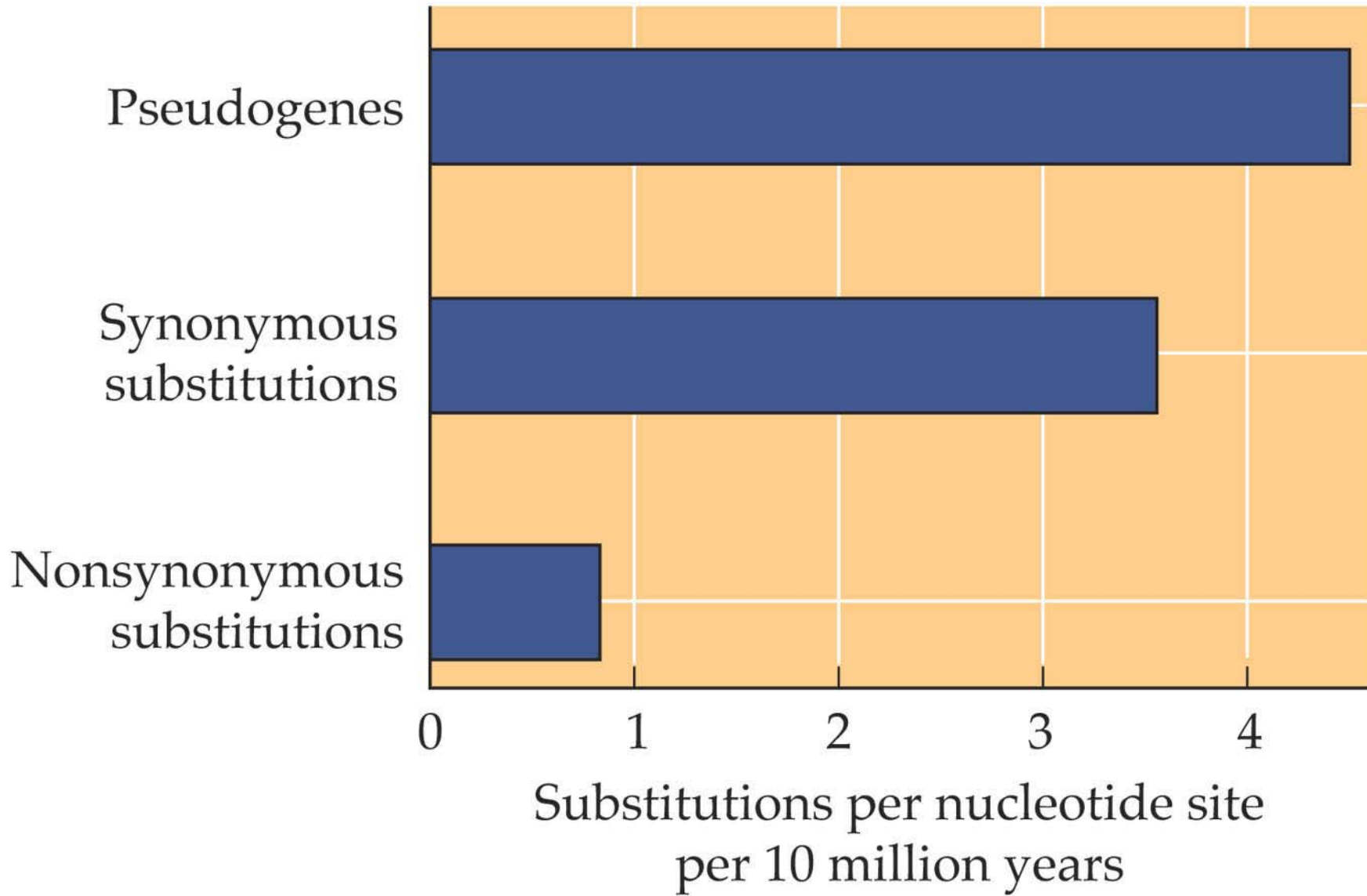
Figure 26.1 *When One Base Does or Doesn't Make a Difference*



Genetic Code (DNA → amino acid)

		Second Letter				
		T	C	A	G	
First Letter	T	TTT } Phe TTC } TTA } Leu TTG }	TCT } TCC } Ser TCA } TCG }	TAT } Tyr TAC } TAA } Stop TAG } Stop	TGT } Cys TGC } TGA } Stop TGG } Trp	T C A G
	C	CTT } CTC } Leu CTA } CTG }	CCT } CCC } Pro CCA } CCG }	CAT } His CAC } CAA } Gln CAG }	CGT } CGC } Arg CGA } CGG }	T C A G
	A	ATT } ATC } Ile ATA } ATG } Met	ACT } ACC } Thr ACA } ACG }	AAT } Asn AAC } AAA } Lys AAG }	AGT } Ser AGC } AGA } Arg AGG }	T C A G
	G	GTT } GTC } Val GTA } GTG }	GCT } GCC } Ala GCA } GCG }	GAT } Asp GAC } GAA } Glu GAG }	GGT } GGC } Gly GGA } GGG }	T C A G

Figure 26.3 Rates of Base Substitution Differ



The Evolution of Genome Size

- Several rounds of duplication and mutation may lead to formation of a **gene family**, a group of **homologous genes with related functions**.
- There is evidence that the globin gene family arose by gene duplication.

= Neofunctionalization

- To estimate the time of the first globin gene duplication, a **gene tree** can be created.
- Based on the gene tree, the two globin gene clusters are estimated to have split about 450 mya.

Figure 26.9 A Globin Family Gene Tree

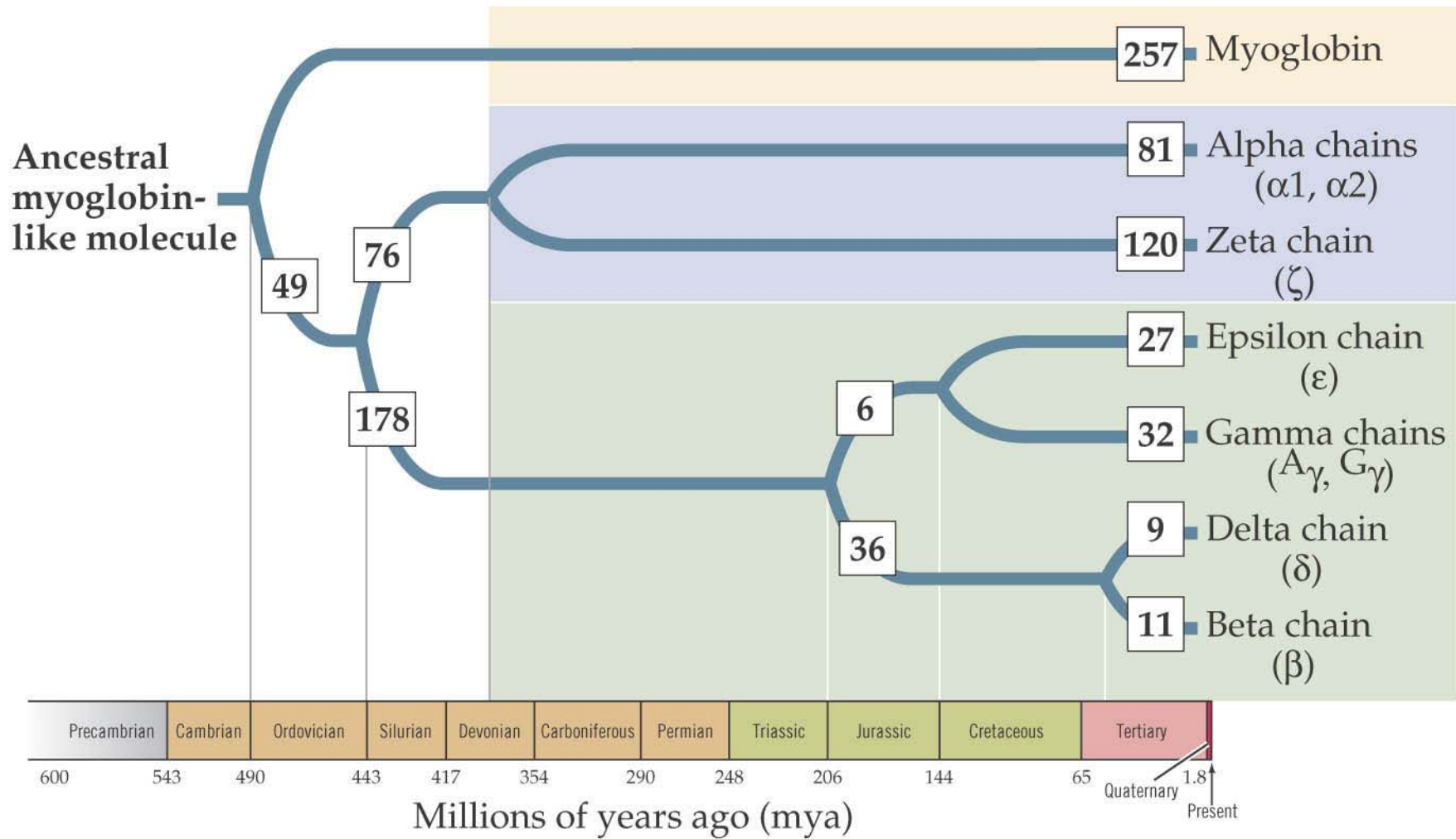
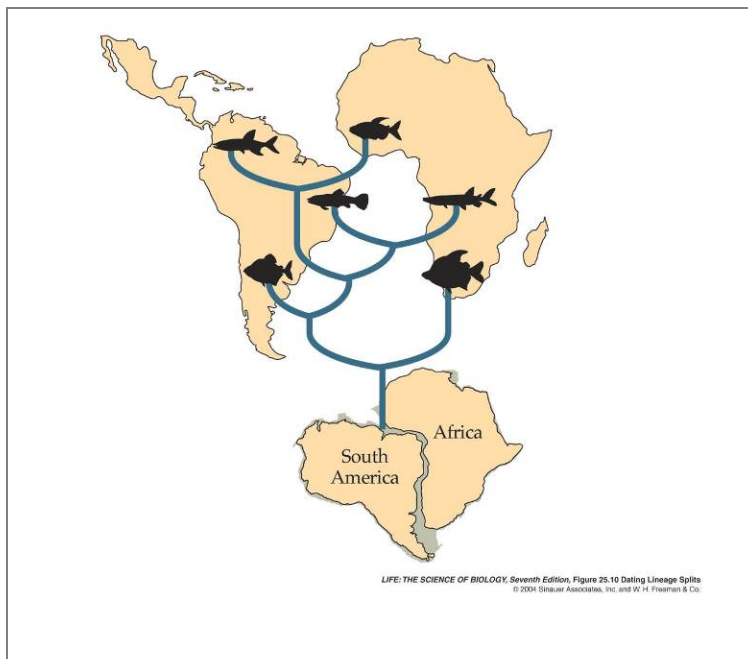
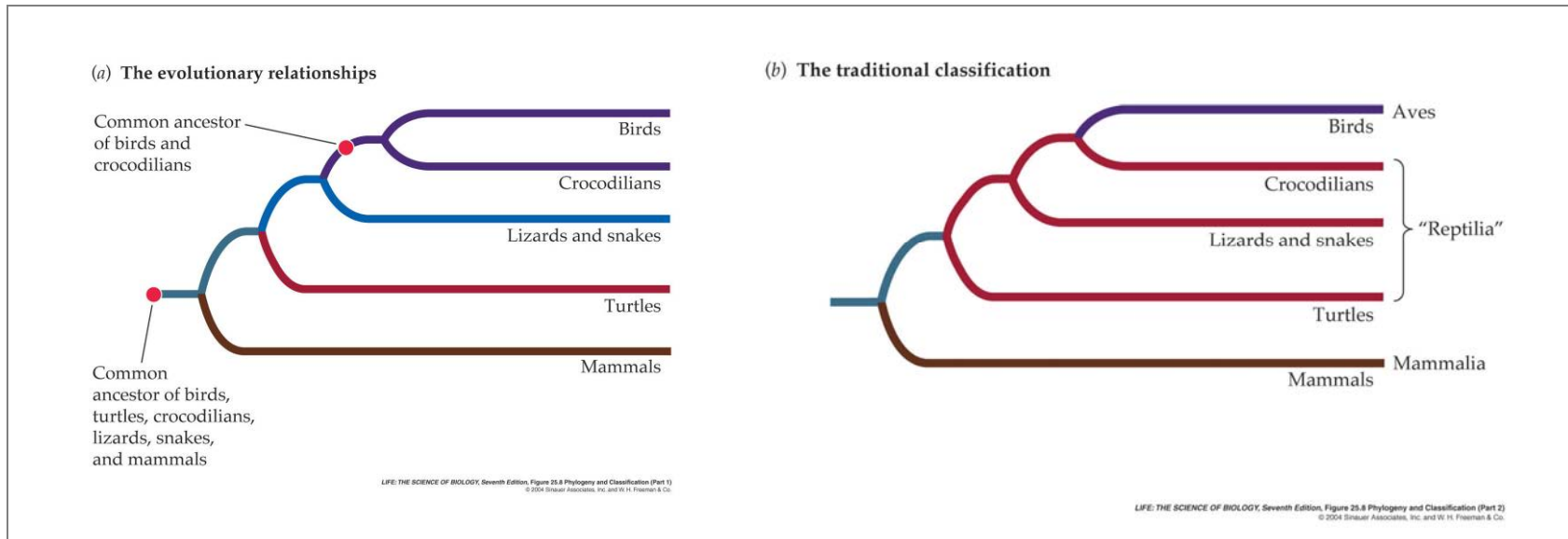


Figure 25.8 *Phylogeny and Classification*



What are the major biological principles illustrated in these two examples?